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148478

From: Yu, Misook
Sent: Tuesday, March 22, 2005 11:07 AM
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Subject: 09/830,338

Pls do interference search of SEQ ID NO:1.

Examiner Misook Yu, Ph.D.
571-272-0839 (Phone)
Art Unit 1642
REM-3A18 (Room)
REM-3C18 (Mail Box)

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Searcher: _____
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Date Searcher Picked up: _____
Date Completed: _____
Searcher Prep/Rev. Time: _____
Online Time: _____

Type of Search

NA#: _____ AA#: _____
Interference: _____ SPDI: _____
S/L: _____ Oligomer: _____
Encode/Transl: _____
Structure#: _____ Text: _____
Inventor: _____ Litigation: _____

Vendors and cost where applicable

STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other(Specify): _____

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OM protein - protein search, using sw model

Run on: March 26, 2005, 09:08:03 ; Search time 27 Seconds
(without alignments)
3878.987 Million cell updates/sec

Title: US-09-830-338-1
Perfect score: 7308
Sequence: 1 MATQOKASDERISQFDHNL.....SKYLTIQKWLFPSPFIQK 1403

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents AA:*
1: /cgn2.6/prodata/1/iaa/5A COMB.pcp.*
2: /cgn2.6/prodata/1/iaa/5B COMB.pcp.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	7308	100.0	1403	US-09-705-872-1	Sequence 1, Appli
2	6691	91.6	1295	US-09-705-872-3	Sequence 3, Appli
3	6373.5	87.2	1232	US-08-836-134-2	Sequence 2, Appli
4	6373.5	87.2	1232	US-09-493-784-2	Sequence 2, Appli
5	5955	81.5	1151	US-08-836-134-23	Sequence 23, Appli
6	5955	81.5	1151	US-09-493-784-23	Sequence 23, Appli
7	462	6.3	618	US-08-511-485-8	Sequence 8, Appli
8	462	6.3	618	US-09-212-971-8	Sequence 8, Appli
9	462	6.3	618	US-08-800-929A-8	Sequence 8, Appli
10	462	6.3	618	US-08-569-749-2	Sequence 2, Appli
11	462	6.3	618	US-09-617-053A-8	Sequence 8, Appli
12	462	6.3	618	US-09-069-023-29	Sequence 29, Appli
13	462	6.3	618	US-09-201-936-8	Sequence 8, Appli
14	462	6.3	618	US-09-011-356-8	Sequence 8, Appli
15	462	6.3	618	US-09-672-717-223	Sequence 223, App
16	462	6.3	618	US-09-201-933-8	Sequence 8, Appli
17	462	6.3	618	US-09-689-366-2	Sequence 2, Appli
18	462	6.3	618	PCT-US96-12860-2	Sequence 2, Appli
19	457.5	6.3	612	US-09-212-971-14	Sequence 14, Appli
20	457.5	6.3	612	US-08-800-929A-14	Sequence 14, Appli
21	457.5	6.3	612	US-08-569-749-14	Sequence 14, Appli
22	457.5	6.3	612	US-09-617-053A-14	Sequence 14, Appli
23	457.5	6.3	612	US-09-689-366-14	Sequence 14, Appli
24	457.5	6.3	612	PCT-US96-12860-14	Sequence 14, Appli
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26	441.5	6.0	591	US-09-011-356-42	Sequence 42, Appli
27	441.5	6.0	591	US-09-672-717-229	Sequence 229, Appl

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37	437.5	6.0	496	4	US-09-011-356-10	Sequence 10, Appli
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44	436.5	6.0	613	4	US-09-949-016-10878	Sequence 10878, A
45	435	6.0	1140	4	US-09-579-692B-8	Sequence 8, Appli

ALIGNMENTS

RESULT 1
US-09-705-872-1
; Sequence 1, Application US/09705872
; Patent No. 6617429
; GENERAL INFORMATION:
; APPLICANT: Joh-E IKEDA
; APPLICANT: Kenji YAMAMOTO
; TITLE OF INVENTION: APOPTOSIS INHIBITORY PROTEIN, GENE ENCODING THE PROTEIN
; FILE OF INVENTION: AND CDNA THEREOF
; FILE REFERENCE: 2000-1110/LC/00653
; CURRENT APPLICATION NUMBER: US/09/705,872
; CURRENT FILING DATE: 2000-11-06
; PRIOR APPLICATION NUMBER: 09/239,797
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 1
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-705-872-1

Query Match	100.0%;	Score 7308;	DB 4;	Length 1403;
Best Local Similarity	100.0%;	Pred. No. 0;		
Matches 1403;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
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DB	121	HDCCGFLNKDVGNTAKYDIRVKNLKSRLGKMYQEEEARLASFRNWPFFYQGISPCV	180	
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DB	181	LSEAGFVFTGKQDTVQCSCGCLGNWEGDDPWKEHAKWFPKCEFLRKSSEETQYI	240	
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DB	301	KAGLFTGTGKDIQVQCSGCLGKQEGDDPDDHTRCFNCPNCFLLQNMKSSAEVTPDLOS	360	

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RESULT 2
US-09-705-872-3
; Sequence 3, Application US/09705872
; Patent No. 6617429
; GENERAL INFORMATION:
; APPLICANT: Joh-E IKEDA
; APPLICANT: Kenji YAMAMOTO
; TITLE OF INVENTION: APOPTOSIS INHIBITORY PROTEIN, GENE ENCODING THE PROTEIN
; TITLE OF INVENTION: AND CDNA THEREOF
; FILE REFERENCE: 2000-1110/LC/00653
; CURRENT APPLICATION NUMBER: US/09705,872
; CURRENT FILING DATE: 2000-11-06
; PRIOR APPLICATION NUMBER: 09/239,797
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: Patent in Ver. 2.0
; SEQ ID NO 3
; LENGTH: 1295
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-705-872-3

Query Match 91.6%; Score 6691; DB 4; Length 1295;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 1282; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

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QY 661 FDDVAVFKSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDLAEAGVDED 720
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Db	121	HPDCGFLNKDVGNIAKYDIRVKULKSRLRGGKRWYQEBEARLASFRNWPYVQGISPCV	180
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Db	181	LSEAGFYFTGKODTVQCFSCGGCLGNWEEGDPMKEHAKWPKCEFLRSKKSSSEITQYI	240
Qy	241	QSYKGFVDITGEHFPVNSVQRELPMSAYCNDISIFAYEELRLDSFKDWPRESAVGVAALA	300
Db	241	QSYKGFVDITGEHFPVNSVQRELPMSAYCNDISIFAYEELRLDSFKDWPRESAVGVAALA	300
Qy	301	KAGLFYTGITKDIIQCFSCGGCLEKWQEGDDPLDDHTRCFPNCPPLOQNKSSAEVTPDLOS	360
Db	301	KAGLFYTGITKDIIQCFSCGGCLEKWQEGDDPLDDHTRCFPNCPPLOQNKSSAEVTPDLOS	360
Qy	361	RGEICELETTSSNLESDSITAVGPIVEMAQGAQWFOEAKNLNEOLRAAYTSASFRHMS	420
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Db	421	LLDISSLATDHLIGCDLSITASKHISKPQOEPLVLPVFGNLNSVMCEGASGKTVLL	480
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Db	601	APFPYNTVCILRKLFSHNMTLRKFVMYFGKNOSLOKI QKTPLFVAACIAHWFQYPPDPS	660
Qy	661	FDDVAVPKSYMERLSLNKATABILKATVSSCGELALKGFFSCFEPNDDDLAEAGVDED	720
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Qy	841	MQLLRGLWQICPOAYFSWVSEHLLVLALKTAYQSNVTVAACSPFVLQFQRTLTLGALNL	900
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Qy	961	ERNLAEKEDNVKSYMDQORASPDLSGYWKLSPKQYKIPCLVEDVNDIDVQGDMLLEIL	1020
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DB 1150 SLPNFISKLINLEGOQPPDETSKFAVILGSLNLEBELILPTGDIYRVAKLIIOQCQ 1209
QY 1261 QLHCLRVLSFFKTLNDDSVVEI 1282
DB 1210 QLHCLRVLSFFKTLNDDSVVEI 1231

RESULT 4
US-09-493-784-2
; Sequence 2, Application US/09493784
; Patent No. 6429011
; GENERAL INFORMATION:
; APPLICANT: Mackenzie, Alex E.
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mahadevan, Mani S.
; APPLICANT: McLean, Michael
; APPLICANT: Roy, Natalie
; APPLICANT: Ikeda, Joh-e
; TITLE OF INVENTION: Neuronal Apoptosis Inhibitor Protein, Gene Sequence and
; Patent No. 6429011
; TITLE OF INVENTION: Mutations Causative of Spinal Muscular Atrophy
; FILE REFERENCE: 3477-112, 033477/139914
; CURRENT APPLICATION NUMBER: US/09/493,784
; CURRENT FILING DATE: 2000-01-28
; PRIOR APPLICATION NUMBER: 08/836,134
; PRIOR FILING DATE: 1997-06-20
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 2
; LENGTH: 1232
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-493-784-2

Query Match 87.2%; Score 6373.5; DB 4; Length 1232;
Best Local Similarity 95.9%; Pred. No. 0;
Matches 1229; Conservative 1; Mismatches 1; Indels 51; Gaps 1;
QY 1 MATQKASDRISQFDHNLPELSALLGLDAVOLAKELBEEBOKERAKWKQGYNSQMRSE 60
DB 1 MATQKASDRISQFDHNLPELSALLGLDAVOLAKELBEEBOKERAKWKQGYNSQMRSE 60
QY 61 AKELKTFVTYEPYSSWIPQMAAAGFYFTGVKSGIOCFCCSLILFAGLTRLPIEDHKRF 120
DB 61 AKELKTFVTYEPYSSWIPQMAAAGFYFTGVKSGIOCFCCSLILFAGLTRLPIEDHKRF 120
QY 121 HPDGGFLNKGDNVGNIAKYDIRVNLKSLRGLRGKRYQOEERARLASFRNWPYVQGISPCV 180
DB 121 HPDGGFLNKGDNVGNIAKYDIRVNLKSLRGLRGKRYQOEERARLASFRNWPYVQGISPCV 180
QY 181 LSEAGFVTKQDTVQCFSCGGLGNWEEGDDPWKSHAKFPKCEFLRSKKSSEETIYI 240
DB 181 LSEAGFVTKQDTVQCFSCGGLGNWEEGDDPWKSHAKFPKCEFLRSKKSSEETIYI 240
QY 241 QSYKGFVDITGEHFVNSWQRELPMASAYCNDISIFAYEELRLDSFKDWPRESAVGVAALA 300
DB 241 QSYKGFVDITGEHFVNSWQRELPMASAYCNDISIFAYEELRLDSFKDWPRESAVGVAALA 300
QY 301 KAGLFYTGKIDIVQCFSCGGLKQWQEGDDPLDDHTRCFPNCPFLQNMKGSAAETPDLOS 360
DB 301 KAGLFYTGKIDIVQCFSCGGLKQWQEGDDPLDDHTRCFPNCPFLQNMKGSAAETPDLOS 360
QY 361 RGECELLETTSSENLEDSTAVGPIVPEMAQGAQWFOEAKNLEQLRAAYTSASFRHMS 420
DB 361 RGECELLETTSSENLEDSTAVGPIVPEMAQGAQWFOEAKNLEQLRAAYTSASFRHMS 420

QY 421 LLDISSDLATDHLGCDLSIASKHISKPOEPLVLPVFGNLNSVMCVGEAGSGKTVLL 480
DB 421 LLDISSDLATDHLGCDLSIASKHISKPOEPLVLPVFGNLNSVMCVGEAGSGKTVLL 480
QY 481 KKIAPLWASGCCPLNRFQVLYLSLSTSRPDEGLASII CDOLLEKEGSVTMCVRNIIO 540
DB 481 KKIAPLWASGCCPLNRFQVLYLSLSTSRPDEGLASII CDOLLEKEGSVTMCVRNIIO 540
QY 541 QLNQVLFLLDDYKEICSIPOVIGKLIQKNHLSRTCLLIIVRTNRARDIRYLETILEIK 600
DB 541 QLNQVLFLLDDYKEICSIPOVIGKLIQKNHLSRTCLLIIVRTNRARDIRYLETILEIQ 600
QY 601 AFFPNTVTCILRLKLFSHNMTLRKFMVYFGNQSLQIKTQKTPLFVAACAHWFQVFPDPS 660
DB 601 AFFPNTVTCILRLKLFSHNMTLRKFMVYFGNQSLQIKTQKTPLFVAACAHWFQVFPDPS 660
QY 661 FDDVAVFKSYMERLSLRNKATATILKATVSSCGELALKGFFSCCFEENDDDLAAGVDED 720
DB 661 FDDVAVFKSYMERLSLRNKATATILKATVSSCGELALKGFFSCCFEENDDDLAAGVDED 720
QY 721 EDLTWCLMSKFTAQRLRPFYFLSPAFOEFLAGMRLIELLSDRQEHQDLGLYHLKQINS 780
DB 721 EDLTWCLMSKFTAQRLRPFYFLSPAFOEFLAGMRLIELLSDRQEHQDLGLYHLKQINS 780
QY 781 PMMTVSAYNNFLNYSVSLPSTKAGPKIVSHLLHLVDNKESENSENDDYLKHQPEISIQ 840
DB 781 PMMTVSAYNNFLNYSVSLPSTKAGPKIVSHLLHLVDNKESENSENDDYLKHQPEISIQ 840
QY 841 MOLLRLGLWQICPOAYFMSVSEHLLVLAUKTAYQSNVAAACSPFVLQFGRTLTILGALNL 900
DB 841 MOLLRLGLWQICPOAYFMSVSEHLLVLAUKTAYQSNVAAACSPFVLQFGRTLTILGALNL 900
QY 901 QYFFDHPESLSLLRSIHPIRGNKTSIPRAHFSVLETCFCKSQVPTIDODYASAFEPNNEW 960
DB 901 QYFFDHPESLSLLRSIHPIRGNKTSIPRAHFSVLETCFCKSQVPTIDODYASAFEPNNEW 960
QY 961 ERNLAEKEDNVKSYNDMQRASPDLSGTGYWKLSPKQYKIPCLVDVNDIDVVGQDMLIL 1020
DB 961 ERNLAEKEDNVKSYNDMQRASPDLSGTGYWKLSPKQYKIPCLVDVNDIDVVGQDMLIL 1020
QY 1021 MTVPASORIELHLNHSRGFTESIRPALELSKASVTKCSISKLELSAAEQELLLTLPSE 1080
DB 1021 MTVPASORIELHLNHSRGFTESIRPALELSKASVTKCSISKLELSAAEQELLLTLPSE 1080
QY 1081 SLEVSSTIQSQDIQIFPNLDKFLCKELSVLEGNINVSFVPIPEEPFNHMEKLIQISA 1140
DB 1081 SLEVSSTIQSQDIQIFPNLDKFLCKELSVLEGNINVSFVPIPEEPFNHMEKLIQISA 1140
QY 1141 EYDPSKLVKLIQNSPNLHVFLKCNFFSDFGSLMTLWVSKKLTETIKFSDSFFQVAPFVA 1200
DB 1141 EYDPSKLVKLIQNSPNLHVFLKCNFFSDFGSLMTLWVSKKLTETIKFSDSFFQVAPFVA 1149
QY 1201 SLPNFISKLINLEGOQPPDETSKFAVILGSLNLEBELILPTGDIYRVAKLIIOQCQ 1260
DB 1150 SLPNFISKLINLEGOQPPDETSKFAVILGSLNLEBELILPTGDIYRVAKLIIOQCQ 1209
QY 1261 QLHCLRVLSFFKTLNDDSVVEI 1282
DB 1210 QLHCLRVLSFFKTLNDDSVVEI 1231

RESULT 5
US-08-836-134-23
; Sequence 23, Application US/08836134A
; Patent No. 6020127
; GENERAL INFORMATION:
; APPLICANT: Mackenzie, Alex E.
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mahadevan, Mani S.
; APPLICANT: McLean, Michael
; APPLICANT: Roy, Natalie
; APPLICANT: Ikeda, Joh-e
; TITLE OF INVENTION: Neuronal Apoptosis Inhibitor Protein, Gene Sequence and

```

; Patent No. 6020127
; TITLE OF INVENTION: Mutations Causative of Spinal Muscular Atrophy
; FILE REFERENCE: 3477-112, 033477/139914
; CURRENT APPLICATION NUMBER: US/08/836,134A
; CURRENT FILING DATE: 1997-06-20
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 23
; LENGTH: 1151
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-08-836-134-23

Query Match      81.5%; Score 5955; DB 3; Length 1151;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 1139; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 MATQOKASDERISQPDHNLPELSALLGLDVAQLAKELEEEERAKQKGYNSQMRSE 60
DB 10 MATQOKASDERISQPDHNLPELSALLGLDVAQLAKELEEEERAKQKGYNSQMRSE 69
QY 61 AKRLKTFVTYEPYSSWIPQEMAAGFYFTGVKSGIQCFCCSLILFAGLTRLPIEDHKRF 120
DB 70 AKRLKTFVTYEPYSSWIPQEMAAGFYFTGVKSGIQCFCCSLILFAGLTRLPIEDHKRF 129
QY 121 HPDCGFLNKDVGNIAKYDIRVKNLKSRLRGKMYQEEEARLASFRNPPFVVOGISPCV 180
DB 130 HPDCGFLNKDVGNIAKYDIRVKNLKSRLRGKMYQEEEARLASFRNPPFVVOGISPCV 189
QY 181 LSEAGFVFTGKQDTVQCFCGCGCLNWEEDGDPWKEHAKWPKCFEFLRSKSSSEITQYI 240
DB 190 LSEAGFVFTGKQDTVQCFCGCGCLNWEEDGDPWKEHAKWPKCFEFLRSKSSSEITQYI 249
QY 241 QSYKGFVDITGSHFVNSWQRELPMASAYCNDISIFAYEELRLDPSFKDWPRESAVGVAALA 300
DB 250 QSYKGFVDITGSHFVNSWQRELPMASAYCNDISIFAYEELRLDPSFKDWPRESAVGVAALA 309
QY 301 KAGLFTYGTIKDVIQVFCGCGCLKEQWEGDDPLDDHTRCPNCPFLQNMKSSAEVTPDLQS 360
DB 310 KAGLFTYGTIKDVIQVFCGCGCLKEQWEGDDPLDDHTRCPNCPFLQNMKSSAEVTPDLQS 369
QY 361 RGECELLETTSESNLDSIAVGPVPEMAQGEAQWFOEAKNLQRLAAVTSASFRHMS 420
DB 370 RGECELLETTSESNLDSIAVGPVPEMAQGEAQWFOEAKNLQRLAAVTSASFRHMS 429
QY 421 LLDISSDLATDHLGCDLSIAKSHISKPVQEPVLVPEVFGNLNSVNCVEGEAGSGKTULL 480
DB 430 LLDISSDLATDHLGCDLSIAKSHISKPVQEPVLVPEVFGNLNSVNCVEGEAGSGKTULL 489
QY 481 KKIAFLWASGCCPLNRFOLVFLYLSLSSTRPDEGLASIIICDQLLEKSGSVTEMCVRNIIQ 540
DB 490 KKIAFLWASGCCPLNRFOLVFLYLSLSSTRPDEGLASIIICDQLLEKSGSVTEMCVRNIIQ 549
QY 541 QLKNOVLFLDDYKICSPQVIGKLIQKNHLSRTCLLIATVNRDRIIRYLETILEIK 600
DB 550 QLKNOVLFLDDYKICSPQVIGKLIQKNHLSRTCLLIATVNRDRIIRYLETILEIQ 609
QY 601 APPFYNTVTCILRKLSHNNTRLRKMYVFGKNSQLOKIQKTPLFVAATCAHWFQFPDPBS 660
DB 610 APPFYNTVTCILRKLSHNNTRLRKMYVFGKNSQLOKIQKTPLFVAATCAHWFQFPDPBS 669
QY 661 FDDVAVFYSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDLAEAGVDED 720
DB 670 FDDVAVFYSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDLAEAGVDED 729
QY 721 EDLTWCLMSKPTAQRLLRPYRFLSPAQEFLLAGMRLIELLSDSDRQHQDLGLYHLKQINS 780
DB 730 EDLTWCLMSKPTAQRLLRPYRFLSPAQEFLLAGMRLIELLSDSDRQHQDLGLYHLKQINS 789
QY 781 PMMTVSAYNNFLNYVSSLPSTVAGPKIVSHLLHLVDNKESLENI SENDDYLKHQPEISLIQ 840
DB 790 PMMTVSAYNNFLNYVSSLPSTVAGPKIVSHLLHLVDNKESLENI SENDDYLKHQPEISLIQ 849

RESULT 6
US-09-493-784-23
; Sequence 23, Application US/09493784
; Patent No. 6429011
; GENERAL INFORMATION:
; APPLICANT: Mackenzie, Alex E.
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mahadevan, Mani S.
; APPLICANT: McLean, Michael
; APPLICANT: Roy, Natalie
; APPLICANT: Ikeda, John-e
; TITLE OF INVENTION: Neuronal Apoptosis Inhibitor Protein, Gene Sequence and
; Patent No. 6429011
; TITLE OF INVENTION: Mutations Causative of Spinal Muscular Atrophy
; FILE REFERENCE: 3477-112, 033477/139914
; CURRENT APPLICATION NUMBER: US/09/493,784
; CURRENT FILING DATE: 2000-01-28
; PRIOR APPLICATION NUMBER: 08/836,134
; PRIOR FILING DATE: 1997-06-20
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 23
; LENGTH: 1151
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-493-784-23

Query Match      81.5%; Score 5955; DB 4; Length 1151;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 1139; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 841 MQLLRGLMQICPQAYFSVMVSEHLLVLAKTAYQSNVTAAACSPFVLQFLOGRTLTLGALML 900
DB 850 MQLLRGLMQICPQAYFSVMVSEHLLVLAKTAYQSNVTAAACSPFVLQFLOGRTLTLGALML 909
QY 901 QYFFDHPESLSLLRSIHPIRGNKTSPPRAHFSVLETCFDKSOVPTTDQDYASAFEPNNEW 960
DB 910 QYFFDHPESLSLLRSIHPIRGNKTSPPRAHFSVLETCFDKSOVPTTDQDYASAFEPNNEW 969
QY 961 ERNLAEKEDNVKSYMDMORRASPDLSGTGYWKLSPKQYKIPCLLEVVDNDIDVVGQDMLEIL 1020
DB 970 ERNLAEKEDNVKSYMDMORRASPDLSGTGYWKLSPKQYKIPCLLEVVDNDIDVVGQDMLEIL 1029
QY 1021 MTFVSASQRIELHNLHNSRGFIESIRPALELSKASVTKCSISKLELSAAQELLLTLPSE 1080
DB 1030 MTFVSASQRIELHNLHNSRGFIESIRPALELSKASVTKCSISKLELSAAQELLLTLPSE 1089
QY 1081 SLEVSGTTQSQDQIPFNLDKFLCLKELSVLDLGNINVSFVPEEPNPFHMKLLIQISA 1140
DB 1090 SLEVSGTTQSQDQIPFNLDKFLCLKELSVLDLGNINVSFVPEEPNPFHMKLLIQISA 1149
QY 1141 E 1141
DB 1150 E 1150
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Db 250 QSYKGFVDITGEHFNVSQVRELPMASAYCNDISFAYEELRLDSDFKDWPRESAVGVAALA 309
Qy 301 KAGLFYTGKIDVQPCSCGCLKEKQEGDDPLDDHTRCPNCPFPLOKMKSSAEVTPDLQS 360
Db 310 KAGLFYTGKIDVQPCSCGCLKEKQEGDDPLDDHTRCPNCPFPLOKMKSSAEVTPDLQS 369
Qy 361 RGECELLETTSESNELESIAGVPIVPEMAQGEAOWFOEAKNLEOLRAAYTSASFRMS 420
Db 370 RGECELLETTSESNELESIAGVPIVPEMAQGEAOWFOEAKNLEOLRAAYTSASFRMS 429
Qy 421 LLDISSDLATDLHLLGCDLSIASKHISKVPQEPVLVPEVFGNLSNVMCVEGEAGSKTVLL 480
Db 430 LLDISSDLATDLHLLGCDLSIASKHISKVPQEPVLVPEVFGNLSNVMCVEGEAGSKTVLL 489
Qy 481 KKIAPFLWASGCCPLNRFOLVFLYSLSSTRPDEGLASIIICDQLEKEGVTMCNRIIQ 540
Db 490 KKIAPFLWASGCCPLNRFOLVFLYSLSSTRPDEGLASIIICDQLEKEGVTMCNRIIQ 549
Qy 541 QLKNOVLFLDDYKEICSIPOVIGKLIQKHLISRTCLLIAVTRNARDIRRYLEILEIK 600
Db 550 QLKNOVLFLDDYKEICSIPOVIGKLIQKHLISRTCLLIAVTRNARDIRRYLEILEIQ 609
Qy 601 APPFYNTVCILRLKLSHNNTRLKFMVYFGKQSLQIKQKTLFVAACIAHWFQYPPFDS 660
Db 610 APPFYNTVCILRLKLSHNNTRLKFMVYFGKQSLQIKQKTLFVAACIAHWFQYPPFDS 669
Qy 661 FDDVAVFKSYMERLSLRNKATAILKATVSSCGELALKGFFSCCFEFDNDDLAEGVDBD 720
Db 670 FDDVAVFKSYMERLSLRNKATAILKATVSSCGELALKGFFSCCFEFDNDDLAEGVDBD 729
Qy 721 EDLTMCMSKFTAQRLLRPYRFLSPAFQFLAGMLRIELLDSDRHOQDLGLYHLKQINS 780
Db 730 EDLTMCMSKFTAQRLLRPYRFLSPAFQFLAGMLRIELLDSDRHOQDLGLYHLKQINS 789
Qy 781 PMWTVSAYNNFLNYSSLSPTKAGPKIVSHLLHLVDNKSLENISENDYLLKHQPEISLQ 840
Db 790 PMWTVSAYNNFLNYSSLSPTKAGPKIVSHLLHLVDNKSLENISENDYLLKHQPEISLQ 849
Qy 841 MQLLRGLMWCICQAFYSVMSEHLLVLALKTAYQSNVTAAACSPFVLQFLQGRITLTIGALNL 900
Db 850 MQLLRGLMWCICQAFYSVMSEHLLVLALKTAYQSNVTAAACSPFVLQFLQGRITLTIGALNL 909
Qy 901 QYFFDHPESLSLRSIHPIRGKTSIPRAHFSVLETCFDKSOVPTIDQDYASAFEPMEW 960
Db 910 QYFFDHPESLSLRSIHPIRGKTSIPRAHFSVLETCFDKSOVPTIDQDYASAFEPMEW 969
Qy 961 ERNLAEKEDNVKSYMDMQRRASPDLSGTGYWKLSPKQYKIPCLEVDVNDIDVVQDMLEIL 1020
Db 970 ERNLAEKEDNVKSYMDMQRRASPDLSGTGYWKLSPKQYKIPCLEVDVNDIDVVQDMLEIL 1029
Qy 1021 MTVFSASQRIELHLSRGIRESIRPALELSKASVTKCSISKLELSAAEQEILLTLPISLE 1080
Db 1030 MTVFSASQRIELHLSRGIRESIRPALELSKASVTKCSISKLELSAAEQEILLTLPISLE 1089
Qy 1081 SLEVSSTIQSDQIPNLKFKLCKELSDVDLEGNINVSFVPIPEEPFNHMEKLIQISA 1140
Db 1090 SLEVSSTIQSDQIPNLKFKLCKELSDVDLEGNINVSFVPIPEEPFNHMEKLIQISA 1149
Qy 1141 E 1141
Db 1150 E 1150

RESULT 7

US-08-511-485-8
; Sequence 8, Application US/08511485
; Patent No. 5919912
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mackenzie, Alexander E.
; APPLICANT: Baird, Stephen
; TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,

TITLE OF INVENTION: PROBES, AND DETECTION METHODS

NUMBER OF SEQUENCES: 38
CORESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/511,485
FILING DATE: 04-AUG-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 07540/002001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 618 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: both
MOLECULE TYPE: protein
US-08-511-485-8

Query Match 6.3%; Score 462; DB 2; Length 618;

Best Local Similarity 24.7%; Pred. No. 5.9e-33;

Matches 159; Conservative 93; Mismatches 238; Indels 154; Gaps 24;

Qy 38 LEEBQKRAKQKQKYNQMRSEAKLTFVYEPYSSWIP---QEMAAAGFYFTGVKSG 94
Db 28 LSDWTNSNKKQKMFSCFCE-----LYRMTSTYTFPAGVPSERSLARAGFYITGVNDK 80
Qy 95 IQCFCCSLILFCAGITRLPIEDHKFHPDCGF---LLNKDVGNIAKYDIRVKN----- 144
Db 81 VKCFCCGLMLDNWKLGDSPIQKHKLQYPCSPFIQNLVSLGSKTSKNTSPMNSFAHSLS 140
Qy 145 -----LKSRL-----RGKMY--QEEEARLASFRNWFYVQG 175
Db 141 PTLHSSLSFGSYSLPPNPLNSRAVEDISSRTPNYPYSAMSTEEARFLTYHWP--LTF 198
Qy 176 ISPCVLSAGFYFTCKQDTVQCFCGCGCLGNWEEGDDPWKEHAKWPKCEFLRSKKSSE 235
Db 199 LSPSELARAGFYITGGRVACFACGGLKSNWEPKDDAMSEHRRHFPNCPFL---ENSE 255
Qy 236 ITQYTSYKGFVDITGEHFNVSQVRELPMASAYCNDISFAYEELRLDSDFKDWPRESAYG 295
Db 256 TLRFSIS-----NLSMQI-----HAARMRTFMYWPSVPVQ 286
Qy 296 VAALAKAGFYTGKIDVQPCSCGCLKEKQEGDDPLDDHTRCPNCPFPLOKMKSSAEVT 355
Db 287 PEQLASAGFYVGRNDDVKFCGDCGLRCWESGDDPWVEHAKWFPCEFLIRMGQ--EFV 345
Qy 356 PDQSR--GELCELLETTSESNELESIAGVPIVPEMAQGEAOWFOEAKNLEOLRAAYTSA 414
Db 346 DEIQRYPHLLQLLSTSDTTGEE--ADPPIIHFGPGESS--SEDVMMNTFVVSALAM 402
Qy 415 SF-----RHMSLLDISD-----ATDLGLGCDLSIAS 442
Db 403 GENRDLVKQTVLSKITTTGENYKVTNDIVSALLNAEDEKREBEKEKQAEEMASDDLILR 462
Qy 443 KHISKPQVQPLVLPVEFQNL---NSVMCVEGAGSGKTVL-----LKKIAFLWASG----- 490

Db 463 KRMALFOQLTCLVLPILNLLKANVINKQEHDIKQKTOIPIQARELIDTIWVKGNAAAN 522
QY 491 ----CCPLNRFQVPLVYLSSTR-----PDGLASIIICDQLEK--EGSVTEMCMRNII 539
Db 523 IFKNC---LKEIDSTLYKNLFVDKNMKYIPTEDVSGLSLEEQRLRQERTCKVCMDKEV 579
QY 540 QOLKNQVFLDDYKICSPQVIGKLIQKNHLSRTCLLIIVRT 583
Db 580 ----SVVPIPCGHLVVC---QECAPSLRKCPCICRGIKGTVRT 615
RESULT 8
US-09-212-971-8
; Sequence 8, Application US/09212971B
; Patent No. 6107041
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: Mackenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Teang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF IAPS AND
; TITLE OF INVENTION: NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; FILE REFERENCE: 07891/009002
; CURRENT APPLICATION NUMBER: US/09/212,971B
; EARLIER FILING DATE: 1998-12-16
; EARLIER APPLICATION NUMBER: 60/017,354
; EARLIER FILING DATE: 1996-04-26
; EARLIER APPLICATION NUMBER: 60/030,590
; EARLIER FILING DATE: 1996-11-14
; EARLIER APPLICATION NUMBER: 08/800,929
; EARLIER FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-212-971-8
Query Match 6.3%; Score 462; DB 3; Length 618;
Best Local Similarity 24.7%; Pred. No. 5.9e-33;
Matches 159; Conservative 93; Mismatches 238; Indels 154; Gaps 24;
QY 38 LEEBQKRAKQKQVNSQMRSEAKRLTFVYFYSSWIP---QEMAAAGFYFTGVKSG 94
Db 28 LSDWTNSNKKQKDYFSCB-----LYRMSTYSTFPAGVPVVSERSLARAGFYTTGVNDK 80
QY 95 IOCFCCSLIFCAGLTRLPIDHKKRPHDPDGP---LLNKKDVGNIAKYDIRVKN----- 144
Db 81 VKFCFCCGLMDLNNKLGDSPIQHKQLYFSCFIONLVASLSTGSKNTSPMRNSFAHLS 140
QY 145 -----LKSRL-----RGCKMRY--QEEBARLASFRNWPFFYVQG 175
Db 141 PYLEHSSLFSGSYSSLPNPLNSRAVEDISSRTTPYSYAMSTEARPLTYIHWMP--LTF 198
QY 176 ISPCVLSEAGVFTGKQDVTQFCSCGCGCLGNWEGDDPWKEHAKWPPKCFPLRSKXSEE 235
Db 199 LSPSELARAGFYIIGDVRVACFACGKLSNWEKDDAMSEHRRHFPNCPFL---ENSLE 255
QY 236 ITQYIQSYKGFVDITCEHPVNSWQRELPMASACNDSIFAYEELRLDSFKDWPRESAVG 295
Db 256 TURFSIS-----NLMSQT-----HAARMRTFMTWPSVPVQ 286
QY 296 VAALAKAGLYTGIKIDIVQCFSCGCGLEKQWQEGDDPLDDHTRCFPNCPFLQNMKSSAEVT 355
Db 287 PEQLASAGFYVGRNDDVKFCGDCGLRCWESGDDPWVEHAKWPPKCFPLRMKQ--EFV 345
QY 356 PLQSR--GELCBLLTTSNLSIEDSIAGVPIPVEMAQQAQWFOBAKNLINEOLRAAYTSA 414
Db 346 DEIQGRYPHLLQLLSTSDTTGEEN--ADPPIIHFGPGESS--SEDVAVMNTFVVKSALEM 402

QY 415 SF-----RHMSLDDISSDI-----ATDHLGCDLSIAS 442
Db 403 GFNRDLVKQTVLSKITLTGENTYKTVNDIVSALLNAEDKEKEEKEKQAEEMASDDLSLR 462
QY 443 KHISKVQBPQLVLPVFGNL---NSVMCVGBAGSGKTVL-----LKKIAFLWASG----- 490
Db 463 KRMALFOQLTCLVLPILNLLKANVINKQEHDIKQKTOIPIQARELIDTIWVKGNAAAN 522
QY 491 ----CCPLNRFQVPLVYLSSTR-----PDGLASIIICDQLEK--EGSVTEMCMRNII 539
Db 523 IFKNC---LKEIDSTLYKNLFVDKNMKYIPTEDVSGLSLEEQRLRQERTCKVCMDKEV 579
QY 540 QOLKNQVFLDDYKICSPQVIGKLIQKNHLSRTCLLIIVRT 583
Db 580 ----SVVPIPCGHLVVC---QECAPSLRKCPCICRGIKGTVRT 615

RESULT 9

US-08-800-929A-8
; Sequence 8, Application US/08800929A
; Patent No. 6133437
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G
; APPLICANT: Mackenzie, Alexander E
; APPLICANT: Liston, Peter
; APPLICANT: Baird, Stephen
; APPLICANT: Teang, Benjamin K
; APPLICANT: Pratt, Christine
; TITLE OF INVENTION: DETECTION AND MODULATION OF
; TITLE OF INVENTION: IAPS AND NAIP FOR THE DIAGNOSIS AND TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: DISEASE
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Clark & Elbing LLP
; STREET: 176 Federal Street
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/800,929A
; FILING DATE: 13-FEB-1997
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 60/030,590
; FILING DATE: 14-NOV-1996
; APPLICATION NUMBER: 60/017,354
; FILING DATE: 26-APR-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Bieker-Brady, Kristina
; REGISTRATION NUMBER:
; REFERENCE/DOCKET NUMBER: 07891/009001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-428-0200
; TELEFAX: 617-428-7045
; TELEX:
; INFORMATION FOR SEQ ID NO: 8:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 618 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-800-929A-8

Query Match 6.3%; Score 462; DB 3; Length 618;
Best Local Similarity 24.7%; Pred. No. 5.9e-33;


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; TITLE OP INVENTION: DISEASE
; FILE REFERENCE: 07891/009003
; CURRENT APPLICATION NUMBER: US/09/617,053A
; CURRENT FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 08/800,929
; PRIOR FILING DATE: 1997-02-13
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-617-053A-8

Query Match      6.3%; Score 462; DB 3; Length 618;
Best Local Similarity 24.7%; Pred.No.5.e-33;
Matches 159; Conservative 93; Mismatches 238; Indels 154; Gaps 24;

Qy   38 LEEBQEKERAKMQKGYNSOMSEAKRLKFTFYYPYSWIP---QEWAAAGFYPTGVKSG 94
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   28 LSDWTNSENKQMKYDFSC_-----LYRMTYSTFTFAGVPVSRSLARAGFYTTGVNDK 80

Qy   95 IQCFCCSILLFAGLTRLPIEDKHFRPHDCGF--LLNKDVGNTAKYDIRVN-----144
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   81 VKCFCCGLMLDNWKLGDSPIQKHQLYPCSFIOQLVSASLGSTSKNTSPMRNSFAHLS 140

Qy   145 -----LKSRL-----RGGMRY--OEBAERLASPNWPFFVQG 175
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   141 PTLEHSILFSGYSSSLPPNLNSRAVEDISSRRNPFSYAMSTEAFPLTHMP--LTF 198

Qy   176 ISPCVLSEAGFVTGKQDTVOCFSCGGCLGNWERGDDPWKEHAKWFPKCEFLRSKKSEE 235
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   199 LSPSELARAGPYIGPDRVACFACGGKLSNWEKCDAMSERRHFNCPEL---ENSLE 255

Qy   236 ITQYIQQYGFVDITGEHFVNSVQRELPMASAYCNDSIFAYELRLDSFKDWPRESAVG 295
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   256 TLRFSIS-----NLSMGT-----HAARMRTFWYFPSSVPVQ 286

Qy   296 VAALAKAGLFYGIKDIVQCFSCGGCLEKWQEGDDPLDDHTRCFPNPCFLQNWKSSAEVT 355
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   287 PEQLASAGFYVGRNDDVKFCGCGGLRWESGDDPWVEHAKWFPREFLRMGQ-EFV 345

Qy   356 PDLOQR-GEICELLETTSESNLEDSIAVGPIPVMAOGEAQWFOEAKNLEQLRAAYTSA 414
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   346 DEIQGRYPHLLEQLLSTDTTGEEN--ADPPIIHFGPGESS-SEDVMMNTPVVKSALEM 402

Qy   415 SF-----RHMSLLDISDL-----ATDHLLGCDSLIAS 442
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   403 GFNRDLVKQTVLISKLTITGENYKTNTDIVSALLNAEDKEEBEKEQAEEASMDLSLR 462

Qy   443 KHISKPVQEPVLVLPVEVRNL--NSVMCVEGEAGSGKTVL----LKIAFIAMWSG-----490
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   463 KRNWALFQOLTCTVLPILDNLKANVINKQEHDIIKQKTQIPLQARELIDTIWVGNAAN 522

Qy   491 ----CCPLLARFQLVFVLSLSSTR-----PDEGLASIIICDQLBEK--EGSVTECMERNII 539
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   523 IFRKC---LKEIDSTLYKNLVFDKNNKYIPTEDVSYGLSLEEQRLQBERTCKVCMDKEV 579

Qy   540 QQLKNQVLFLLDDYKEICSIPOVIGKUIQKNHLSRTCLLIART 583
       | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db   580 -----SVVFIPCQHLVWC---QECAPSLRKCPICRGIKNGTVT 615
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RESULT 12
US-09-069-023-29
; Sequence 29, Application US/09089023A
; Patent No. 6348573
; GENERAL INFORMATION:
; APPLICANT: Nunez, Gabriel
; APPLICANT: Inohara, Naohiro
; APPLICANT: Koseki, Takeyoshi
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR IDENTIFYING APOPTOSIS
; TITLE OF INVENTION: SIGNALING PATHWAY INHIBITORS AND ACTIVATORS

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; FILE REFERENCE: UM-03333
; CURRENT APPLICATION NUMBER: US/09/069,023A
; CURRENT FILING DATE: 1998-04-27
; NUMBER OF SEQ ID NOS: 38
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 29
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-069-023-29

Query Match      6.3%; Score 462; DB 3; Length 618;
Best Local Similarity 26.0%; Pred.No.5.9e-33;
Matches 155; Conservative 82; Mismatches 214; Indels 146; Gaps 22

Qy 38 LEEBEQKERAKMKGYNYSOMRSEAKRLKTFVYYPYSSWIP--QEMAAAGFYFTGVKSG 94
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 28 LSDWTNNGKQMKDYFSC-----LYRMTYSTFPAGVPVSRSLARAGFYITGVNDK 80

Qy 95 IQCFCCSLILFAGLTLPIEDHKRPHDCGF---LLNKOVGNTAKYDIRVKN----- 144
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 81 VKFCFCGLMDLNWKJGDSPIQKHQLYPCSCFIQLNLSVASLGSTSKNTSPMNSFAHLS 140

Qy 145 -----LKSRL-----RGKMY--QEEEARLASFRNWFYVQG 175
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 141 PTLHSLFSGSYSSLSPNPLNSRAVEDISSRTNPYSYAMSTBEARFLTTHWP--LTF 198

Qy 176 ISPCVLSEAGFVFTGKODTQCFCSCGCLGNWESGDDPWKEHAKWFPKCEFLRSKSSFE 235
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 199 LSPSELARAGFYIIGPGRVACFACGGKJLNWEPKDDAMSEHRRHFPNCPEL---ENSLE 255

Qy 236 ITQVIQSYKGFVDITGEHFVNSWQRELPMASAYCNDSIFAYEELRLDSFKDWPRESAVG 295
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 256 TLAPSL-----NLSMQT-----HAARWTFMYWPSVPVQ 286

Qy 296 VAALAKAGLYTGKDTIQVCFSCGCGLEKWOEGDDPLDHTRCFPNCPFLQNMKSSAEVT 355
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 287 PEQLASAGFYVGRNDVDKFCDDGLRCWESGDDPWVEHAKWFPKCEFLRMKQG--EFV 345

Qy 356 PDLOSR--GELCELLETTSESNLDSIAGPIVPMAOGAQQFOEAKNLNQLRAAYTSA 414
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 346 DEIQGVPHLEQLLSTSTDTGEEN--ADPPIIHFGGESS--SEDAVMNTFPVKSALEM 402

Qy 415 SFRHMSLLDISDLATDHLGLCDLSIAKSHISKVQEP-LVLPVEFGNLNSVMCEGEGAG 473
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 403 GFNR-----DL-----VKQTVQSKILTTGENYKTVNDIVSALLNAE 438

Qy 474 SGKTVLK-KIAFLWAGCCPLNRFOLVFYLSLSSTRPDGLASIIDQLLEKEGVSYTE 532
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 439 DEKREEKEKQAEASDDLSLRKRNWALFQOLTCVLP-----ILDNLK----- 484

Qy 533 MCMRNITIQLNQVFLFLDDYKEICSTPQVIGKLIQRNHLISRTCLLTAVRTNRARDI 589
      . : : : : : : : : : : : : : : : : : : : : : : : : : : : : : :
Db 485 ---ANVINKQEHDI-----KQKTIPLQARELID-----TILVKGNAANI 523

RESULT 13
US-09-201-936-8
Sequence 8, Application US/09201936
Patent No. 6541457
GENERAL INFORMATION:
APPLICANT: Korneluk, Robert G.
APPLICANT: MacKenzie, Alexander E.
APPLICANT: Baird, Stephen
APPLICANT: Liston, Peter
TITLE OF INVENTION: MAMMALIAN IAP GENE FAMILY, PRIMERS,
TITLE OF INVENTION: PROBES, AND DETECTION METHODS
FILE REFERENCE: 07891/003003
CURRENT APPLICATION NUMBER: US/09/201,936
CURRENT FILING DATE: 1998-12-01
EARLIER APPLICATION NUMBER: 09/011,356
EARLIER FILING DATE: 1998-02-04
EARLIER APPLICATION NUMBER: PCT/IB96/01022

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; EARLIER FILING DATE: 1996-08-05
; EARLIER APPLICATION NUMBER: 08/576,956
; EARLIER FILING DATE: 1995-12-22
; EARLIER APPLICATION NUMBER: 08/511,485
; EARLIER FILING DATE: 1995-08-04
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 8
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-201-936-8

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Query Match	6.38;	Score 462;	DB 4;	Length 618;
Best Local Similarity	24.7%;	Pred. No. 5.9e-33;		
Matches 159;	Conservative	93;	Mismatches 238;	Indels 154;
Gaps	24;			
Qy	38	LEBEEQERAKMQKGYNSOMRSEAKRLKTFVYEPYSWIP--QEMAAAGFYTGKVGSG	94	
Db	28	LSDWTSNKKQMKYDFSC-----LYRMTSTYTFPAGVPVRSRLARAGFYITGVNDK	80	
Qy	95	IQCFCCSLIIFGAGLTLPIEDHKRFHPDCG--LLNKDVGNTAKYDIRVN-----	144	
Db	81	VKCFCCGLMLDNWKLGDSPIQKHQLYPSCFIQNLVSASLGSTSKNTSPMRNSFAHSLS	140	
Qy	145	-----LKSRL-----RGGKMY--QESSEARLASPNRPVFFVQG	175	
Db	141	PTLEHSSLPFSGSYSLPPNPPLNSRAVEDISSRTPNPSYAMSTEARFLTYHWP--LTF	198	
Qy	176	ISPCVLSEAGFVFTGKODTVQCFSCGGLGNWREGDDPWKEHAKWPKCFELRSKKSEE	235	
Db	199	LSPSELARAGFYITGPGDRVACFACGGKLSNWEPKDDAMSHRRHFPNCPFL---ENSLE	255	
Qy	236	ITQYIQSVKGFVDITGEHFVNSWVQRELPMASAYCNDISI PAYEELRLDSPKDPRESAVG	295	
Db	256	TLRPSIS-----NLWQT-----HAARMTFYWVSSVPVQ	286	
Qy	296	VAALAKAGLFYTGIKDIVQCFSCGGCLEKWEQGDPLDDHTRCPNCPFLQNMKSSAEVT	355	
Db	287	PEQLASAGFYVGNRDDVKFCGCGGLRCWESGDDPWVEHAKWFPCEFLLRMRGQ--EFV	345	
Qy	356	PDLOSR--GEICELLETTSESNLDSIAGPITPEMAQGEAQWFOEAKNLNQLRAAYTSA	414	
Db	346	DEIQGRYPHLEQLGLSTDTGGEN--ADPPIIHFGPGESS--SEDVAMVNTPVVKSALEM	402	
Qy	415	SF-----RHMSLLDISDL-----ATDHLILGCDLSTAS	442	
Db	403	GFNRDLVKQTVLSKILITGGENYKTVNDIVSALLNAEDEKREEKEQAEWASDLSLR	462	
Qy	443	KHISKPVQEPILVLEPVGNL---NSVMCVEGEASGKTVL---LKKTAFLWASG-----	490	
Db	463	KNRMALFQQLTCLVPLIDNLKLVNINKQEHDIKQKQIPLQARELIDTTWVGNAAN	522	
Qy	491	----CCPLLNRFLQVFLYLSISSTR-----PDEGLASICTDQLLEK--RGSVTECMRNII	539	
Db	523	IFKNK---LKEIDSTLYXNLFVDNMKVIPTEDVYSGLSLEBQLRLQBERTCKVCMDKEV	579	
Qy	540	QQLKXQVLFLLDDYKICISIPQVTGKLIQKNHLSRTCLLIIVRT	583	
Db	580	-----SVVFIPCGHLVVC---QECAPSLRKCPICRGIIGKTVRT	615	

580 -----SVFIPOGHLVVC---QECAPSLRKPCIRGIIKGTVRT 615
 RESULT 15
 US-09-672-717-223
 - Sequence 223, Application US/09672717
 - Patent No. 6673917
 - GENERAL INFORMATION:
 : APPLICANT: Korneluk, Robert G.
 : APPLICANT: LaCasse, Eric

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; APPLICANT: Baird, Stephen
; APPLICANT: Holcik, Martin
; APPLICANT: Young, Sean
; TITLE OF INVENTION: Antisense IAP Nucleic Acids and Uses
; TITLE OF INVENTION: Thereof
; FILE REFERENCE: 07891/025001
; CURRENT APPLICATION NUMBER: US/09/672,717
; CURRENT FILING DATE: 2000-09-28
; NUMBER OF SEQ ID NOS: 231
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 223
; LENGTH: 618
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-672-717-223

Query Match      6.3%; Score 462; DB 4; Length 618;
Best Local Similarity 24.7%; Pred. No. 5.9e-33;
Matches 159; Conservative 93; Mismatches 238; Indels 154; Gaps 24;

QY 38 LEEEEKERAKQKGYNSQMRSEAKLKTFTVTEPYSSWIP---QEMAAAGFYFTGVKSG 94
Db 28 LSDWTNSNQKKYDFSCB-----LYRMTYSTFPAGVPVSESLARAGFYITGVNDK 80
QY 95 IQCFCCSLILFGAGLTRLPDIEHKRPHPCGP---LLNRDVGNGIAKYDIRVKN----- 144
Db 81 VKCFCCGLMLDNWKLGDSPIQKHQKLYPSCSFQNLVLSASLGSTSKNTSPMENSFAHLS 140
QY 145 -----LKSRL-----RGGKORY--QEEEARLASFRNWPFFVQG 175
Db 141 PTLHSSLSFGSYSLPPNPLNSRAVEDISSRTNFPYSAMSTEEARFLTTHWP--LTF 198
QY 176 ISPCVLSEAGFVFTGQDVTQCFSCGCGCIKNNEEGDPWKEHAKWPKCFELRSKXSEE 235
Db 199 LSPSELARAGFYIIGPDGVACGCGKUSNWEPPKDDAMSEHRRHFPNCFEL---ENSLE 255
QY 236 ITQYIQSYKGFVDITGEHFVNSVQRELPMASAYCNDISIFAYEELRLDSFKDWPRESAVG 295
Db 256 TLRFSIS-----NLSMQT-----HAARMRTFTWPSVPVQ 286
QY 296 VAALAKAGLYTGIKDIVQCFSCGCGLEKQWQGGDDPLDHTRCFPNCPFLONKXSAEVT 355
Db 287 PEQLASAGFYVYGRNDDVKCFGCGGLRCWESGDDPWVEHAKWPFPRCEFLIRMKQ-BFV 345
QY 356 PLQSR-GECELLETTSNLEDSTAVGPIVPEMAQEAQWFOEAKNLEQIRAAYTSA 414
Db 346 DEIQRYPHLEQLLSTSDTTGEEN--ADPPIIHFGPGESS-SEDAVMNTPVVKSALEM 402
QY 415 SF-----RHMSLLDISDL-----ATDHLGCDLSIAS 442
Db 403 GNRDLVKQTVLSKILTTGENTKYTVNDIVSALLNAEDEKREKEKEQAEMASDDLILIR 462
QY 443 KHISKPQPLVLPVEFGNL---NSVMCEGAGSGKTVL-----LKKIAPLWASG----- 490
Db 463 KNRMALFQQLTCVLPILDNLKANVINKQEHDI IKQKTOIPLQARELIDTIWVGNAAN 522
QY 491 -----CCPLNRQLVFLYLSSTR-----PDGLASIIDQLLEK--EGSVTEMCMRNII 539
Db 523 IFKNC---LKEIDSTLYKNLFDVKNMKYIPTEDVSGLSLEEQRLRLQERTCKVCMDREV 579
QY 540 QQLKNQVFLDDDYKEICSIPOVIGKLIQKNHLSRTCLLIAVRT 583
Db 580 -----SVFIPGHLVVC---QECAPSLRKCPICRGIINGTVRT 615
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Job time : 30 secs

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: March 26, 2005, 09:12:04 ; Search time 101 Seconds
(without alignments)
4599.352 Million cell updates/sec

Title: US-09-830-338-1
Perfect score: 7308
Sequence: 1 MATOQKASDERISQFDHLL.....SKYLTIQKWLPPSPFIQK 1403

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1407402 seqs, 331100923 residues
Total number of hits satisfying chosen parameters: 1407402

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA:*

- 1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep.*
- 2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep.*
- 3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep.*
- 4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep.*
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- 6: /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pep.*
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- 10: /cgn2_6/ptodata/1/pubpaa/US09B_PUBCOMB.pep.*
- 11: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep.*
- 12: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pep.*
- 13: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep.*
- 14: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep.*
- 15: /cgn2_6/ptodata/1/pubpaa/US10C_PUBCOMB.pep.*
- 16: /cgn2_6/ptodata/1/pubpaa/US10D_PUBCOMB.pep.*
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- 18: /cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep.*
- 19: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep.*
- 20: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	7308	100.0	1403	8	US-08-913-322-22
2	7308	100.0	1403	8	US-08-913-322-24
3	7308	100.0	1403	14	US-10-285-408-1
4	3970.5	54.3	782	9	US-09-841-739-9
5	3970.5	54.3	782	14	US-10-449-315-9
6	1975	27.0	385	14	US-10-029-386-33707
7	1903	26.0	898	9	US-09-841-739-11
8	1903	26.0	898	14	US-10-449-315-11
9	1037	14.2	203	14	US-10-029-386-33933
10	661	9.0	1204	9	US-09-841-739-5
11	661	9.0	1204	14	US-10-449-315-5
12	654	8.9	1024	14	US-10-156-733-2
13	654	8.9	1070	14	US-10-221-097-49

14	653	8.9	1024	9	US-09-841-739-2	Sequence 2, Appli
15	653	8.9	1024	14	US-10-449-315-2	Sequence 2, Appli
16	641	8.8	1024	9	US-09-864-921-97	Sequence 97, Appl
17	641	8.8	1024	17	US-10-766-682-97	Sequence 97, Appl
18	545.5	7.5	118	9	US-09-925-299-1033	Sequence 1033, Ap
19	545.5	7.5	118	10	US-09-925-299-1033	Sequence 1033, Ap
20	536.5	7.3	738	14	US-10-221-097-48	Sequence 48, Appl
21	462	6.3	618	9	US-09-974-592-8	Sequence 8, Appli
22	462	6.3	618	9	US-09-201-936-8	Sequence 8, Appli
23	462	6.3	618	14	US-10-153-668-338	Sequence 338, App
24	462	6.3	618	14	US-10-207-655-200	Sequence 200, App
25	462	6.3	618	14	US-10-232-286-2	Sequence 2, Appli
26	462	6.3	618	15	US-10-366-307-4	Sequence 4, Appli
27	462	6.3	618	15	US-10-361-270-3	Sequence 3, Appli
28	462	6.3	618	15	US-10-260-708-63	Sequence 63, Appl
29	462	6.3	618	16	US-10-636-065-223	Sequence 223, App
30	462	6.3	618	16	US-10-600-272-8	Sequence 8, Appli
31	462	6.3	618	16	US-10-730-476A-78	Sequence 78, Appl
32	462	6.3	618	17	US-10-934-717-2	Sequence 2, Appli
33	457.5	6.3	612	9	US-09-974-592-14	Sequence 14, Appl
34	457.5	6.3	612	14	US-10-232-286-14	Sequence 14, Appl
35	457.5	6.3	612	17	US-10-934-717-14	Sequence 14, Appl
36	441.5	6.0	591	9	US-09-201-936-42	Sequence 42, Appl
37	441.5	6.0	591	16	US-10-636-065-229	Sequence 229, App
38	441.5	6.0	591	16	US-10-600-272-42	Sequence 42, Appl
39	440	6.0	600	9	US-09-974-592-12	Sequence 12, Appl
40	437.5	6.0	496	9	US-09-974-592-10	Sequence 10, Appl
41	437.5	6.0	496	9	US-09-201-936-10	Sequence 10, Appl
42	437.5	6.0	496	16	US-10-636-065-225	Sequence 225, App
43	437.5	6.0	496	16	US-10-600-272-10	Sequence 10, Appl
44	436.5	6.0	604	14	US-10-232-286-4	Sequence 4, Appli
45	436.5	6.0	604	14	US-10-141-618-6	Sequence 6, Appli

ALIGNMENTS

RESULT 1
US-08-913-322-22
; Sequence 22, Application US/08913322
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mackenzie, Alexander E.
; APPLICANT: Roy, Natalie
; APPLICANT: Robertson, George
; APPLICANT: Tamai, Katsu
; TITLE OF INVENTION: USER OF NEURONAL APOPTOSIS INHIBITOR
; TITLE OF INVENTION: (NAIP)
; FILE REFERENCE: 07891/013001
; CURRENT APPLICATION NUMBER: US/08/913,322
; CURRENT FILING DATE: 1997-09-12
; EARLIER APPLICATION NUMBER: PCT/IB97/00142
; EARLIER FILING DATE: 1997-01-17
; EARLIER APPLICATION NUMBER: GB 9601108.5
; EARLIER FILING DATE: 1996-01-19
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 22
; LENGTH: 1403
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-913-322-22

Query Match 100.0%; Score 7308; DB 8; Length 1403;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 MATOQKASDERISQFDHLLPELSALLGLDVAQLAKELEEESEKERAKMKQKNSQMRSE 60
DB 1 MATOQKASDERISQFDHLLPELSALLGLDVAQLAKELEEESEKERAKMKQKNSQMRSE 60
QY 61 AKRLKTFVTEPYSSWIPQEMAAAGFYTGKVGKQCFCSCSLILFGAGLTRLPIDHKRF 120

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Db 1141 EYDPSKLVKLIQNSPNLHVPHLKC�FFSDFGSLMTMLVSCCKLTETKIFSDSFFQAVPFVA 1200
Qy 1201 SLPNFISLKIINLEGOQPPDEETSEKPAYILGSLNLEELILPTGDTGYRVAKLIIQOCQ 1260
Db 1201 SLPNFISLKIINLEGOQPPDEETSEKPAYILGSLNLEELILPTGDTGYRVAKLIIQOCQ 1260
Qy 1261 QHCLRVLSPPKTLNDDSVVEIAKVAISGGFQKLENKLSINHKITBEGYRNFQALDNM 1320
Db 1261 QHCLRVLSFFKTLNDDSVVEIAKVAISGGFQKLENKLSINHKITBEGYRNFQALDNM 1320
Qy 1321 PNLQELDISRHFTCEIKAQATTVKSLSQCVLRRLRLRLNMLSWLLDADDIALLNVKMR 1380
Db 1321 PNLQELDISRHFTCEIKAQATTVKSLSQCVLRRLRLRLNMLSWLLDADDIALLNVKMR 1380
Qy 1381 HPQSKYLTILOKWILPSPPIIQ 1403
Db 1381 HPQSKYLTILOKWILPSPPIIQ 1403

RESULT 2
US-08-913-322-24
; Sequence 24, Application US/08913322
; Publication No. US20020137028A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mackenzie, Alexander E.
; APPLICANT: Roy, Natalie
; APPLICANT: Robertson, George
; APPLICANT: Tamai, Katsuo
; TITLE OF INVENTION: USER OF NEURONAL APOPTOSIS INHIBITOR
; TITLE OF INVENTION: (NAIP)
; FILE REFERENCE: 07891/013001
; CURRENT APPLICATION NUMBER: US/08/913,322
; CURRENT FILING DATE: 1997-09-12
; EARLIER APPLICATION NUMBER: PCT/IB97/00142
; EARLIER FILING DATE: 1997-01-17
; EARLIER APPLICATION NUMBER: GB 9601108.5
; EARLIER FILING DATE: 1996-01-19
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 24
; LENGTH: 1403
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-913-322-24

Query Match 100.0%; Score 7308; DB 8; Length 1403;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MATQOKASDERISQFDHNLPELSALLGLDVAQLAKELEEEQKERAKQKGYNSQMRSE 60
Db 1 MATQOKASDERISQFDHNLPELSALLGLDVAQLAKELEEEQKERAKQKGYNSQMRSE 60
Qy 61 AKRLKTFVTYEPYSSWIPQEMAAAGFYFTGVKSGIQCFCCSLLIFGAGLTRLPIDHKRF 120
Db 61 AKRLKTFVTYEPYSSWIPQEMAAAGFYFTGVKSGIQCFCCSLLIFGAGLTRLPIDHKRF 120
Qy 121 HPDCGFLNKKOVNTAKYDIRVKNLKSRLRGKMYQEEEARLASFRNPPFYVQGISPCV 180
Db -121 HPDCGFLNKKOVNTAKYDIRVKNLKSRLRGKMYQEEEARLASFRNPPFYVQGISPCV 180
Qy 181 LSEAGFVFTGKQDTVQCFCGCGCLNWEEDGDPWKEHAKWPKCFKLSKSSSEITQYI 240
Db 181 LSEAGFVFTGKQDTVQCFCGCGCLNWEEDGDPWKEHAKWPKCFKLSKSSSEITQYI 240
Qy 241 QSYKGFVDITGHEFVNSWVQRELPMASAYCNDISIYABELRLDPSKDPRESAVGVAALA 300
Db 241 QSYKGFVDITGHEFVNSWVQRELPMASAYCNDISIYABELRLDPSKDPRESAVGVAALA 300
Qy 301 KAGLFYTGIDKIVQCFCGCGCLEKQEGDDPLDDHTRCFPPNCFPQNMKSSAEVTPDLQS 360
Db 301 KAGLFYTGIDKIVQCFCGCGCLEKQEGDDPLDDHTRCFPPNCFPQNMKSSAEVTPDLQS 360
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Db 1141 EYDPSKLVKLIQNSPNLHVPHLKC�FFSDFGSLMTMLVSCCKLTETKIFSDSFFQAVPFVA 1200
Qy 1201 SLPNFISLKIINLEGOQPPDEETSEKPAYILGSLNLEELILPTGDTGYRVAKLIIQOCQ 1260
Db 1201 SLPNFISLKIINLEGOQPPDEETSEKPAYILGSLNLEELILPTGDTGYRVAKLIIQOCQ 1260
Qy 1261 QHCLRVLSPPKTLNDDSVVEIAKVAISGGFQKLENKLSINHKITBEGYRNFQALDNM 1320
Db 1261 QHCLRVLSFFKTLNDDSVVEIAKVAISGGFQKLENKLSINHKITBEGYRNFQALDNM 1320
Qy 1321 PNLQELDISRHFTCEIKAQATTVKSLSQCVLRRLRLRLNMLSWLLDADDIALLNVKMR 1380
Db 1321 PNLQELDISRHFTCEIKAQATTVKSLSQCVLRRLRLRLNMLSWLLDADDIALLNVKMR 1380
Qy 1381 HPQSKYLTILOKWILPSPPIIQ 1403
Db 1381 HPQSKYLTILOKWILPSPPIIQ 1403

RESULT 2
US-08-913-322-24
; Sequence 24, Application US/08913322
; Publication No. US20020137028A1
; GENERAL INFORMATION:
; APPLICANT: Korneluk, Robert G.
; APPLICANT: Mackenzie, Alexander E.
; APPLICANT: Roy, Natalie
; APPLICANT: Robertson, George
; APPLICANT: Tamai, Katsuo
; TITLE OF INVENTION: USER OF NEURONAL APOPTOSIS INHIBITOR
; TITLE OF INVENTION: (NAIP)
; FILE REFERENCE: 07891/013001
; CURRENT APPLICATION NUMBER: US/08/913,322
; CURRENT FILING DATE: 1997-09-12
; EARLIER APPLICATION NUMBER: PCT/IB97/00142
; EARLIER FILING DATE: 1997-01-17
; EARLIER APPLICATION NUMBER: GB 9601108.5
; EARLIER FILING DATE: 1996-01-19
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 24
; LENGTH: 1403
; TYPE: PRT
; ORGANISM: Homo sapiens
US-08-913-322-24

Query Match 100.0%; Score 7308; DB 8; Length 1403;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MATQOKASDERISQFDHNLPELSALLGLDVAQLAKELEEEQKERAKQKGYNSQMRSE 60
Db 1 MATQOKASDERISQFDHNLPELSALLGLDVAQLAKELEEEQKERAKQKGYNSQMRSE 60
Qy 61 AKRLKTFVTYEPYSSWIPQEMAAAGFYFTGVKSGIQCFCCSLLIFGAGLTRLPIDHKRF 120
Db 61 AKRLKTFVTYEPYSSWIPQEMAAAGFYFTGVKSGIQCFCCSLLIFGAGLTRLPIDHKRF 120
Qy 121 HPDCGFLNKKOVNTAKYDIRVKNLKSRLRGKMYQEEEARLASFRNPPFYVQGISPCV 180
Db -121 HPDCGFLNKKOVNTAKYDIRVKNLKSRLRGKMYQEEEARLASFRNPPFYVQGISPCV 180
Qy 181 LSEAGFVFTGKQDTVQCFCGCGCLNWEEDGDPWKEHAKWPKCFKLSKSSSEITQYI 240
Db 181 LSEAGFVFTGKQDTVQCFCGCGCLNWEEDGDPWKEHAKWPKCFKLSKSSSEITQYI 240
Qy 241 QSYKGFVDITGHEFVNSWVQRELPMASAYCNDISIYABELRLDPSKDPRESAVGVAALA 300
Db 241 QSYKGFVDITGHEFVNSWVQRELPMASAYCNDISIYABELRLDPSKDPRESAVGVAALA 300
Qy 301 KAGLFYTGIDKIVQCFCGCGCLEKQEGDDPLDDHTRCFPPNCFPQNMKSSAEVTPDLQS 360
Db 301 KAGLFYTGIDKIVQCFCGCGCLEKQEGDDPLDDHTRCFPPNCFPQNMKSSAEVTPDLQS 360
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QY 361 RGELCELLETTSESNNLEDSIAVGPIVPEMAQGEAOWFOBAKNLNEQLRAAYTSASFRHMS 420
Db 361 RGELCELLETTSESNNLEDSIAVGPIVPEMAQGEAOWFOBAKNLNEQLRAAYTSASFRHMS 420
QY 421 LLDISSDLATDHLGCDLSIAKSHISKPVQEBVLVLPEVFGNLSNVMCVGEAGSGTKVLL 480
Db 421 LLDISSDLATDHLGCDLSIAKSHISKPVQEBVLVLPEVFGNLSNVMCVGEAGSGTKVLL 480
QY 481 KKIAPLWASGCCPLNRRFQVFLVLSLSTRPDEGLASIIICDQLLEKESGVTMCMRNIQ 540
Db 481 KKIAPLWASGCCPLNRRFQVFLVLSLSTRPDEGLASIIICDQLLEKESGVTMCMRNIQ 540
QY 541 QLKNOVLFLDDYKEICSIPOVIGKLIQKNHLSRTCLLIAVTRNRARDIRRYLETILEIK 600
Db 541 QLKNOVLFLDDYKEICSIPOVIGKLIQKNHLSRTCLLIAVTRNRARDIRRYLETILEIK 600
QY 601 APFFYNTVCILRLKLFHNMTLRKFMVYFGKQSLQIKQKTLPLFVAACAHWFQYPPFDS 660
Db 601 APFFYNTVCILRLKLFHNMTLRKFMVYFGKQSLQIKQKTLPLFVAACAHWFQYPPFDS 660
QY 661 FDDVAVFKSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDDDLAEAGVDED 720
Db 661 FDDVAVFKSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDDDLAEAGVDED 720
QY 721 EDLTWCLMSKFTAQRLRPFRFLSPAFOEFLAGMRLLIELLSDRQHQDLGLYHLKQINS 780
Db 721 EDLTWCLMSKFTAQRLRPFRFLSPAFOEFLAGMRLLIELLSDRQHQDLGLYHLKQINS 780
QY 781 PMMTVSAYNNFLNYSLSLSTAGPKIVSHLLHLVDNKESLENISSNDYLLKHQPEISIQ 840
Db 781 PMMTVSAYNNFLNYSLSLSTAGPKIVSHLLHLVDNKESLENISSNDYLLKHQPEISIQ 840
QY 841 MQLRLGLWQICPOAVFSWSEHLVLAKTAVQSNNTVAACSPVLOFLQGRTLTLGALNL 900
Db 841 MQLRLGLWQICPOAVFSWSEHLVLAKTAVQSNNTVAACSPVLOFLQGRTLTLGALNL 900
QY 901 QYFFDHPELSLLRSIHPIRGNTKSPRAHFSVLETCFDKSOVPTTIDQYASAFPMNEW 960
Db 901 QYFFDHPELSLLRSIHPIRGNTKSPRAHFSVLETCFDKSOVPTTIDQYASAFPMNEW 960
QY 961 ERNLAEKNVKSMDMORRASPDISTGYWKLSPQYKIPCLEVDVNDIDVVQDMLEIL 1020
Db 961 ERNLAEKNVKSMDMORRASPDISTGYWKLSPQYKIPCLEVDVNDIDVVQDMLEIL 1020
QY 1021 MTVPESASQRIELHLSHSGFIESIRPALELSKASVTKCSISKLELSAAEQELLLTLPISLE 1080
Db 1021 MTVPESASQRIELHLSHSGFIESIRPALELSKASVTKCSISKLELSAAEQELLLTLPISLE 1080
QY 1081 SLEVSQTIQSQOIFPNLDKFLCKELSVLDLEGNINVSFVPIPEEPFNHMEKLLIQISA 1140
Db 1081 SLEVSQTIQSQOIFPNLDKFLCKELSVLDLEGNINVSFVPIPEEPFNHMEKLLIQISA 1140
QY 1141 EYDPSKLVKLIQNSPNLHVFLHKCNFPDFGSLMTLWLVSKKLTETKFSDFSFFQAVPFVA 1200
Db 1141 EYDPSKLVKLIQNSPNLHVFLHKCNFPDFGSLMTLWLVSKKLTETKFSDFSFFQAVPFVA 1200
QY 1201 SLNPNFISLKIANGEOQFPDEETSEKFAVILGSLNLEHLIILPTGDGIYRAKLIIOQCO 1260
Db 1201 SLNPNFISLKIANGEOQFPDEETSEKFAVILGSLNLEHLIILPTGDGIYRAKLIIOQCO 1260
QY 1261 QLHCLRVLSFFKTLNDDSVVEIAKVAISGGFOKLENKLSINHKITEEGYRNFFOALDNM 1320
Db 1261 QLHCLRVLSFFKTLNDDSVVEIAKVAISGGFOKLENKLSINHKITEEGYRNFFOALDNM 1320
QY 1321 PNLQELDISRHFTPECIKAQATTVKSQCVLRPLRILRLNMLSWLLDADDIALLNVMKER 1380
Db 1321 PNLQELDISRHFTPECIKAQATTVKSQCVLRPLRILRLNMLSWLLDADDIALLNVMKER 1380
QY 1381 HPQSKYLTTLQKWILPFPSPIIQK 1403
Db 1381 HPQSKYLTTLQKWILPFPSPIIQK 1403
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RESULT 3
US-10-285-408-1
; Sequence 1, Application US/10285408
; Publication No. US20030108967A1
; GENERAL INFORMATION:
; APPLICANT: IKEDA, Johe
; TITLE OF INVENTION: Monoclonal Antibodies Against Human Apoptosis Inhibitory Protein 1
; FILE REFERENCE: 2002-1440/WMC/00653
; CURRENT FILING DATE: 2002-11-01
; PRIOR FILING DATE: 2001-04-26
; PRIOR APPLICATION NUMBER: PCT/JPS99/05841
; NUMBER OF SEQ ID NOS: 2
; SEQ ID NO 1
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-285-408-1

Query Match 100.0%; Score 7308; DB 14; Length 1403;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1403; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MATQOKASDERISQPDHNLPELSALLGLDVAOLAKELSEEEQKERAKMKQKGYNSQMRSE 60
Db 1 MATQOKASDERISQPDHNLPELSALLGLDVAOLAKELSEEEQKERAKMKQKGYNSQMRSE 60
QY 61 AKRLKFTVTEYBYSWIPQEMAAAGFYFTGVKSGIQCFCCSLILFCAGLRLPIEDHKRP 120
Db 61 AKRLKFTVTEYBYSWIPQEMAAAGFYFTGVKSGIQCFCCSLILFCAGLRLPIEDHKRP 120
QY 121 HPDCGFLNLKDVGNIAKYDIRVKNLSRLGCKMRYQBEERLASFRNMPFVVGISPCV 180
Db 121 HPDCGFLNLKDVGNIAKYDIRVKNLSRLGCKMRYQBEERLASFRNMPFVVGISPCV 180
QY 181 LSEAGFVTGTQDVTQCFSCGGCLNWEEDDPWKEHAKWPKCFELRSKKSSEETQYI 240
Db 181 LSEAGFVTGTQDVTQCFSCGGCLNWEEDDPWKEHAKWPKCFELRSKKSSEETQYI 240
QY 241 QSYKGFVDITGSHFVNSWQRELPMASVYCNDSIFAYBELRLDSPKDWPRESAVGVAALA 300
Db 241 QSYKGFVDITGSHFVNSWQRELPMASVYCNDSIFAYBELRLDSPKDWPRESAVGVAALA 300
QY 301 KAGLFYTGIDQIVQCFSCGGCLEKMQEGDDPLDDHTRCFPNCPFLQNMKSSAEVTPDLQS 360
Db 301 KAGLFYTGIDQIVQCFSCGGCLEKMQEGDDPLDDHTRCFPNCPFLQNMKSSAEVTPDLQS 360
QY 361 RGELCELLETTSESNNLEDSIAVGPIVPEMAQGEAOWFOBAKNLNEQLRAAYTSASFRHMS 420
Db 361 RGELCELLETTSESNNLEDSIAVGPIVPEMAQGEAOWFOBAKNLNEQLRAAYTSASFRHMS 420
QY 421 LLDISSDLATDHLGCDLSIAKSHISKPVQEBVLVLPEVFGNLSNVMCVGEAGSGTKVLL 480
Db 421 LLDISSDLATDHLGCDLSIAKSHISKPVQEBVLVLPEVFGNLSNVMCVGEAGSGTKVLL 480
QY 481 KKIAPLWASGCCPLNRRFQVFLVLSLSTRPDEGLASIIICDQLLEKESGVTMCMRNIQ 540
Db 481 KKIAPLWASGCCPLNRRFQVFLVLSLSTRPDEGLASIIICDQLLEKESGVTMCMRNIQ 540
QY 541 QLKNOVLFLDDYKEICSIPOVIGKLIQKNHLSRTCLLIAVTRNRARDIRRYLETILEIK 600
Db 541 QLKNOVLFLDDYKEICSIPOVIGKLIQKNHLSRTCLLIAVTRNRARDIRRYLETILEIK 600
QY 601 APFFYNTVCILRLKLFHNMTLRKFMVYFGKQSLQIKQKTLPLFVAACAHWFQYPPFDS 660
Db 601 APFFYNTVCILRLKLFHNMTLRKFMVYFGKQSLQIKQKTLPLFVAACAHWFQYPPFDS 660
QY 661 FDDVAVFKSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDDDLAEAGVDED 720
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Db 661 FDDVAVFKSYMERLSLRNKATAEILKATVSSCGELALKGFFSCCFEFDNDLAEAGVDED 720
QY 721 EDLTWCLMSKFTAQRLRPYRFLSPAFQBLAGMLRIELLDSDRQHQDLGILYHLKQINS 780
Db 721 EDLTWCLMSKFTAQRLRPYRFLSPAFQBLAGMLRIELLDSDRQHQDLGILYHLKQINS 780
QY 781 PMWTVSAYNNFLNYVSSLSSTAGPKIVSHLHLVDNKSLENIENDDYLKHQPEISLQ 840
Db 781 PMWTVSAYNNFLNYVSSLSSTAGPKIVSHLHLVDNKSLENIENDDYLKHQPEISLQ 840
QY 841 MOLLRLGLWQICQAYFSVMSEHLLVLALKTAYQSNVTAAACSPFVLQFLOGRTLTILGALML 900
Db 841 MOLLRLGLWQICQAYFSVMSEHLLVLALKTAYQSNVTAAACSPFVLQFLOGRTLTILGALML 900
QY 901 QYFFDHPELSLRSIHFPKNGKTSPPRAHFSVLETCFDKSPQVPTIDQYASAFPMNEW 960
Db 901 QYFFDHPELSLRSIHFPKNGKTSPPRAHFSVLETCFDKSPQVPTIDQYASAFPMNEW 960
QY 961 ERNLAEKEDNVKSYMDQRRASPDLSGTGWLSPKQYKIPCLEVDVNDIDVVGQDMLEIL 1020
Db 961 ERNLAEKEDNVKSYMDQRRASPDLSGTGWLSPKQYKIPCLEVDVNDIDVVGQDMLEIL 1020
QY 1021 MTVFSASQRIELHLSHRSFGFIESIRPALBELSKASVTKCSISKLELSAAEQELLTLPSLE 1080
Db 1021 MTVFSASQRIELHLSHRSFGFIESIRPALBELSKASVTKCSISKLELSAAEQELLTLPSLE 1080
QY 1081 SLEVSQTIOQDQIPNLKFKCLKLSVDLEGNINVFSPVPEEPNPFHMEKLLIQISA 1140
Db 1081 SLEVSQTIOQDQIPNLKFKCLKLSVDLEGNINVFSPVPEEPNPFHMEKLLIQISA 1140
QY 1141 EYDPSKLVKLIQNSPNLHVFLHKNCFPSDFGSLMTMLVSKKLTETKPSDSFFQAVPFA 1200
Db 1141 EYDPSKLVKLIQNSPNLHVFLHKNCFPSDFGSLMTMLVSKKLTETKPSDSFFQAVPFA 1200
QY 1201 SLPNFISLKLNLGEGQFPDEETSEKFAVILGSLNLELILPTGDIYRVAKLIIQQCQ 1260
Db 1201 SLPNFISLKLNLGEGQFPDEETSEKFAVILGSLNLELILPTGDIYRVAKLIIQQCQ 1260
QY 1261 QLHCLRVLSFFKTLNDDSVVEIAKVAISGGFQKLENLKLINHKITEEGYRNFQALDNM 1320
Db 1261 QLHCLRVLSFFKTLNDDSVVEIAKVAISGGFQKLENLKLINHKITEEGYRNFQALDNM 1320
QY 1321 PNLQELDISRHFTCECAKAQATTVKSLSQCVLRPLRLNMLSWLLDADDIALLNVWKER 1380
Db 1321 PNLQELDISRHFTCECAKAQATTVKSLSQCVLRPLRLNMLSWLLDADDIALLNVWKER 1380
QY 1381 HPQSKYLTILQKWILPSPPIQK 1403
Db 1381 HPQSKYLTILQKWILPSPPIQK 1403

RESULT 4
US-09-841-739-9
; Sequence 9, Application US/09841739
; Patent No. US20020034784A1
; GENERAL INFORMATION:
; APPLICANT: Bertin, John
; TITLE OF INVENTION: NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THERE
; FILE REFERENCE: 07334-329001
; CURRENT APPLICATION NUMBER: US/09/841,739
; CURRENT FILING DATE: 2001-08-29
; PRIOR APPLICATION NUMBER: US 09/697,089
; PRIOR FILING DATE: 2000-10-26
; PRIOR APPLICATION NUMBER: US 60/161,822
; PRIOR FILING DATE: 1999-10-27
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: Fast-Seq for Windows Version 4.0
; SEQ ID NO 9
; LENGTH: 782
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-841-739-9

Query Match 54.3%; Score 3970.5; DB 9; Length 782;
Best Local Similarity 93.6%; Pred. No. 3.5e-308;
Matches 779; Conservative 1; Mismatches 1; Indels 51; Gaps 1;
QY 451 EPLVLPEVFGNLNVMCVGEAGSGKTVLLKKIAFLWASGCCPLLRQLVYFLSLSTR 510
Db 1 EPLVLPEVFGNLNVMCVGEAGSGKTVLLKKIAFLWASGCCPLLRQLVYFLSLSTR 60
QY 511 PDEGLASIIICDOLLEKEGSVTMCMRNIIQOLKNQVFLDDYKEICSIPOVIGKLIQKN 570
Db 61 PDEGLASIIICDOLLEKEGSVTMCMRNIIQOLKNQVFLDDYKEICSIPOVIGKLIQKN 120
QY 571 HLSRTCLLIIVTRNARDIRRYLETILEIKAPFFNTVCILRKLFSHNMTRLRKFMVYFG 630
Db 121 HLSRTCLLIIVTRNARDIRRYLETILEIQAPFFNTVCILRKLFSHNMTRLRKFMVYFG 180
QY 631 KNQSLQKIQTPLFVAALCAHWFOYFPDPSFDDVAVFKSYMERLSLRNKATAEILKATVS 690
Db 181 KNQSLQKIQTPLFVAALCAHWFOYFPDPSFDDVAVFKSYMERLSLRNKATAEILKATVS 240
QY 691 SCGELALKGFFSCCFEFDNDLAEAGVDEDLTWCLMSKETAORLRFYFLSPAFQEF 750
Db 241 SCGELALKGFFSCCFEFDNDLAEAGVDEDLTWCLMSKETAORLRFYFLSPAFQEF 300
QY 751 LAGMRLIELLSDSDRQHQDLGLYHLKQINS PMWTVSAYNNFLNYVSSLSSTKAGPKIVSH 810
Db 301 LAGMRLIELLSDSDRQHQDLGLYHLKQINS PMWTVSAYNNFLNYVSSLSSTKAGPKIVSH 360
QY 811 LLHLVDNKSLENIENDDYLKHQPEISLQMLRLGLWQICQAYFSVMSEHLLVLALKT 870
Db 361 LLHLVDNKSLENIENDDYLKHQPEISLQMLRLGLWQICQAYFSVMSEHLLVLALKT 420
QY 871 AYQSNVTAAACSPFVLQFLOGRTLTIGALNLOVFDHPDESILRSIHFPKNGKTSPPRAH 930
Db 421 AYQSNVTAAACSPFVLQFLOGRTLTIGALNLOVFDHPDESILRSIHFPKNGKTSPPRAH 480
QY 931 FSVLTCFDKSPQVPTIDQYASAFPMNEWERNLAEKEDNVKSYMDMORRASPDLSGTGW 990
Db 481 FSVLTCFDKSPQVPTIDQYASAFPMNEWERNLAEKEDNVKSYMDMORRASPDLSGTGW 540
QY 991 KLSPKQYKIPCLEVDVNDIDVVGQDMLEILMTVFSASQRIELHLSHRSFGFIESIRPAL 1050
Db 541 KLSPKQYKIPCLEVDVNDIDVVGQDMLEILMTVFSASQRIELHLSHRSFGFIESIRPAL 600
QY 1051 SKASVTKCSISKLELSAAEQELLTLPSLESLEVS GTTQSODQIFPNLDKFLCLKELSDV 1110
Db 601 SKASVTKCSISKLELSAAEQELLTLPSLESLEVS GTTQSODQIFPNLDKFLCLKELSDV 660
QY 1111 LEGNINVFSPVPEEPNPFHMEKLLIQISAEYDPSKLVKLIQNSPNLHVFLHKNFFSDF 1170
Db 661 LEGNINVFSPVPEEPNPFHMEKLLIQISAEYDPSKL----- 697
QY 1171 GSLMTMLVSKKLTETKFSDFQAVPFVASFPLNPFISILKLNLEGOQFPDEETSEKFAVI 1230
Db 698 -----VASLPNPFISILKLNLEGOQFPDEETSEKFAVI 729
QY 1231 LGSLSNLELILPTGDIYRVAKLIIQQCQQLHCLRVLSFFKTLNDDSVVEI 1282
Db 730 LGSLSNLELILPTGDIYRVAKLIIQQCQQLHCLRVLSFFKTLNDDSVVEI 781

RESULT 5
US-10-449-315-9
; Sequence 9, Application US/10449315
; Publication No. US20030190679A1
; GENERAL INFORMATION:
; APPLICANT: Bertin, John
; TITLE OF INVENTION: NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THERE
; FILE REFERENCE: 07334-329001
; CURRENT APPLICATION NUMBER: US/10/449,315
; CURRENT FILING DATE: 2003-05-30
; PRIOR APPLICATION NUMBER: US 10/841,739

;; PRIOR FILING DATE: 2001-08-29
;; PRIOR APPLICATION NUMBER: US 09/697,089
;; PRIOR FILING DATE: 2000-10-26
;; PRIOR APPLICATION NUMBER: US 60/161,822
;; PRIOR FILING DATE: 1999-10-27
;; NUMBER OF SEQ ID NOS: 16
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 9
;; LENGTH: 782
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-10-449-315-9

Query Match 54.3%; Score 3970.5; DB 14; Length 782;
Best Local Similarity 93.6%; Pred. No. 3.5e-308;
Matches 779; Conservative 1; Mismatches 1; Indels 51; Gaps 1;

QY	451	EPLVLPEVFGNLNVNVCVEGAGSGKTVLLKIAFLWASGCCPLLNRFLQVFLYLSSTR	510
DB	1	EPLVLPEVFGNLNVNVCVEGAGSGKTVLLKIAFLWASGCCPLLNRFLQVFLYLSSTR	60
QY	511	PDEGLASIIICDQLEKEGVTMCNENIIQOLKNQVLFLLDDYKETSIPQVIGKLIQKN	570
DB	61	PDEGLASIIICDQLEKEGVTMCNENIIQOLKNQVLFLLDDYKETSIPQVIGKLIQKN	120
QY	571	HLSTRCLLIIVRTRNARDIRRYLETILEIKAPFPYNTVCILRLKFSHNNTRLRKFWYFG	630
DB	121	HLSTRCLLIIVRTRNARDIRRYLETILEIQAPFPYNTVCILRLKFSHNNTRLRKFWYFG	180
QY	631	KNQSLQKIQKTPLFVAACIAHMFQYFPDPSDDVAVFKSYMRLSLRNKATAILKATVS	690
DB	181	KNQSLQKIQKTPLFVAACIAHMFQYFPDPSDDVAVFKSYMRLSLRNKATAILKATVS	240
QY	691	SGELALKGFFSCCFNFDDDLAEAGVDEDELTWCLMSKFTAQRLRPVRFISPAQFQF	750
DB	241	SGELALKGFFSCCFNFDDDLAEAGVDEDELTWCLMSKFTAQRLRPVRFISPAQFQF	300
QY	751	LQGMRLIELLSDRQHQDGLYHLKQINSYNNFNLYVSSLPSTKAGPKIVSH	810
DB	301	LQGMRLIELLSDRQHQDGLYHLKQINSYNNFNLYVSSLPSTKAGPKIVSH	360
QY	811	LLHLVDNKSLENISNDYKHQPEISLQMLLRLGLWQICQVAFVSMVSEHLLVLAKT	870
DB	361	LLHLVDNKSLENISNDYKHQPEISLQMLLRLGLWQICQVAFVSMVSEHLLVLAKT	420
QY	871	AYQSNVTAACSPVLOFQGRTLTLGALNLQYFFDHPESLSLLRSIHFSIRGNKTS	930
DB	421	AYQSNVTAACSPVLOFQGRTLTLGALNLQYFFDHPESLSLLRSIHFSIRGNKTS	480
QY	931	FSVLETCPDKSQVPTIDQYASAFEPNNEWERNLAEKEDNVKSYMOMRRASPDLS	990
DB	481	FSVLETCPDKSQVPTIDQYASAFEPNNEWERNLAEKEDNVKSYMOMRRASPDLS	540
QY	991	KLSPKQYKIPCLEVDVNDIDVVGQDMLTILMTVFSASQRIELHNLHNSRGFIESIR	1050
DB	541	KLSPKQYKIPCLEVDVNDIDVVGQDMLTILMTVFSASQRIELHNLHNSRGFIESIR	600
QY	1051	SKASVTKCSISKLELSAAEQELLTLTPSLESLEVSQTIQSDQIIPNLDKFLCLKEL	1110
DB	601	SKASVTKCSISKLELSAAEQELLTLTPSLESLEVSQTIQSDQIIPNLDKFLCLKEL	660
QY	1111	LEGINIVFVPIEPPNFHMEKLLIQISAEDYDPSKLVKLIQNSPNLHVFLKCNFFSD	1170
DB	661	LEGINIVFVPIEPPNFHMEKLLIQISAEDYDPSKLVKLIQNSPNLHVFLKCNFFSD	697
QY	1171	GSIMTLMVSKKLTETIKFSDSPFQVAFVLPNFIKILNLEGQQFPDEETSEKFAYI	1230
DB	698	-----VASLPNFIKILNLEGQQFPDEETSEKFAYI	729
QY	1231	LGSLSNLEELIILPTGDIYRVAKLIIQQCQQLHCLRLVLSFFKTLNDDSVVEI	1282
DB	730	LGSLSNLEELIILPTGDIYRVAKLIIQQCQQLHCLRLVLSFFKTLNDDSVVEI	781

RESULT 6

US-10-029-386-33707
; Sequence 33707, Application US/10029386
; Publication No. US20030194704A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR G
; TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
; FILE REFERENCE: AEOMICA-X-2
; CURRENT APPLICATION NUMBER: US/10/029,386
; NUMBER OF SEQ ID NOS: 34288
; SOFTWARE: Anomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 33707
; LENGTH: 385
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO U80017.1
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 0.91
; OTHER INFORMATION: SWISSPROT HIT: Q13075, EVALUUE 0.00e+00
US-10-029-386-33707

Query Match 27.0%; Score 1975; DB 14; Length 385;
Best Local Similarity 100.0%; Pred. No. 5.2e-149;
Matches 385; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	707	FNDDDLAAGVDEDELTWCLMSKFTAQRLRPVRFISPAQFQFLAGMRLIELLSDRQE	766
DB	1	FNDDDLAAGVDEDELTWCLMSKFTAQRLRPVRFISPAQFQFLAGMRLIELLSDRQE	60
QY	767	HQDLGLYHLKQINSYNNFNLYVSSLPSTKAGPKIVSHLLHLDNKSLENISE	826
DB	61	HQDLGLYHLKQINSYNNFNLYVSSLPSTKAGPKIVSHLLHLDNKSLENISE	120
QY	827	NDLYKHQPEISLQMLLRLGLWQICQVAFVSMVSEHLLVLAKTAYQSNVTAACSPFV	886
DB	121	NDLYKHQPEISLQMLLRLGLWQICQVAFVSMVSEHLLVLAKTAYQSNVTAACSPFV	180
QY	887	FLQGRRTLGLALNLQYFFDHPESLSLLRSIHPIRGNTKSPRAHFSVLETCPDKSQVPTI	946
DB	181	FLQGRRTLGLALNLQYFFDHPESLSLLRSIHPIRGNTKSPRAHFSVLETCPDKSQVPTI	240
QY	947	DQYASAFEPNNEWERNLAEKEDNVKSYMOMRRASPDLSQYKIPKQYKIPCLEVDV	1006
DB	241	DQYASAFEPNNEWERNLAEKEDNVKSYMOMRRASPDLSQYKIPKQYKIPCLEVDV	300
QY	1007	NDIDVVGQDMLTILMTVFSASQRIELHNLHNSRGFTESIRPALELSKASVTKCSISKLELS	1066
DB	301	NDIDVVGQDMLTILMTVFSASQRIELHNLHNSRGFTESIRPALELSKASVTKCSISKLELS	360
QY	1067	RAEQELLTLTPSLESLEVSQTIQSQ 1091	
DB	361	RAEQELLTLTPSLESLEVSQTIQSQ 385	

RESULT 7

US-09-841-739-11
; Sequence 11, Application US/09841739
; Patent No. US20020034784A1
; GENERAL INFORMATION:
; APPLICANT: Bertin, John
; TITLE OF INVENTION: NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THERE
; FILE REFERENCE: 07334-329001
; CURRENT APPLICATION NUMBER: US/09/841,739
; CURRENT FILING DATE: 2001-08-29
; PRIOR APPLICATION NUMBER: US 09/697,089
; PRIOR FILING DATE: 2000-10-26
; PRIOR APPLICATION NUMBER: US 60/161,822
; PRIOR FILING DATE: 1999-10-27

; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 11
; LENGTH: 898
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: majority sequence
US-09-841-739-11

Query Match 26.0%; Score 1903; DB 9; Length 898;
Best Local Similarity 50.9%; Pred. No. 1.2e-142;
Matches 484; Conservative 88; Mismatches 214; Indels 164; Gaps 24;

QY 451 EPLVPEVFGNLSVMCVGEAGSGKTVLLKIAFLWASGCCPLNRRFOLVFLVLSSTR 510
DB 1 EQVLVNGVLGALNSVCIVEGAGSGKSVLLQKIAFLWAGKCKALTQFQVFLVLSSTR 60
QY 511 PDEGLASIIICDQLLEKESGVTMCMRNIIQQLKNQVFLFLDDYKEI----CSIPOVIGKL 566
DB 61 ADGGLASIIICDQLLEKESGVTMCMRNIIQQLKNQVFLFLDDYKEI----CSIPOVIGKL 120
QY 567 IQKNHLSRTCLLIATVNRARDIRRYLETILEIKAPFFYNTVCIILKLFSHNMTRLRKPM 626
DB 121 IQENHLSKTCVLVAVTTERARDIROFGALIAEVGAFEDSAVALLREVLIKEIAELRGLL 180
QY 627 VYFGKNSLQIKTKTFLFVAALCA-HWFQYPPDPSPDDVAVPKSYMERSLRNK-----A 680
DB 181 VQIGKSQSIQNLQKTPFLFVAALCAIQWGESFDSFTDVAVPKSFVDDLILKNKHGVA 240
QY 681 TABILKATVSSCGDLALEGVFHKFDFFELDDVAEAGVDEVDLLTTGLLSKFTAQRLRPY 740
DB 241 AADILKATVSSCGDLALEGVFHKFDFFELDDVAEAGVDEVDLLTTGLLSKFTAQRLKPY 300
QY 741 RFLSPAFQBLAGRLIIELDSDROHQDILGLVHLKQINSMPMTVSAYNNFLNYSSLS 800
DB 301 KFLSKAFQBLAGRLIIELDSDROHQDILGLVHLKQINSMPMTVSAYNNFLNYSSLS 360
QY 801 TKAGPKIVSHLLHLVDNKESLENISENDYLLKHQPEISIQMOLLRLGLWQICQAFYSMS 860
DB 361 VEAGRAVSHLAAVVDNKGSLGLSIADYLLKHQESISIQMOLLQGVKNITQAILSAVS 420
QY 861 EHLVLALKTAVQSNNTAACSPVLQ-FLQGRITLIGALNL-QY---PFDDH-PESLSLR 914
DB 421 INLLVLGITAYQSSITVAALSQVLEAFLOQKSLTLGAGNLPDYLDFDHLFESASALD 480
QY 915 SIHFPIRGNTKSPRAHFSVLETCFDKSVQPTIDQDY--ASAFEPNWEERNIAEKEDNVK 972
DB 481 SIKLSIRGGATASRAKAAVLTGGIDKSEAPTIDETIYPASAVSLFENEQEALATLEVTVK 540
QY 973 SYMDMQRASPDLSGTGYWKLSPQYKIP-----CLEVDVNDIDVV 1012
DB 541 SPSDLNKQAITDGTGFSASSLQIQKRCAGVAGSLVLSLSTCKNIYSLVDASDLTV 600
QY 1013 GDMLEIL--MTVFS---ASORIEHLNHS---RGFTIESIRPALELSKAVTK-----1057
DB 601 GEDHLTIVNLTVLVSHDLASQLEGLTSLGNLKGLELIDALELSEAKLAELG 660
QY 1058 -----CSISKLELSAAEQELLTLTSLSES---LE-----VSGTIQSDQIFPNL 1098
DB 661 KNLKMKCLISLELSAAGEGLLIVKLSLSEPCDLEIQLVSCCLVAGAVQILAILHNL 720
QY 1099 DK--FLCLAKELSDLEGNINVSIVPEEPNPHMEKLIQISAEVDPKSLVKLQNSPN 1156
DB 721 VKLSILDLSLSVLDLGNIAVHSIVPEEPNVLQELTALLLQIGADV-----767
QY 1157 LHVFLKCNFPFSDGLMTLWVSCCKLTIKFSDFSFPQAVFVSLPNFISLKIINLEQ 1216
DB 768 -----GSLSSL-----VASEEIVSLVILGLEQ 791
QY 1217 QPPDETSKFAVI-IGLSLNLFEELILPTGD-----GIYRVAKL-----1254
DB 792 QLTDTBEISILGAFIGLSLNLFEELILAGDWSGDLAFMGVFEVAKLLVFPDFSTKEF 851

QY 1255 -----IIQOQOQLHCLRVLSFPKT-----LNDDSVVEIAKVAISGGFQ 1292
DB 852 LPDPALVQQLSQV--LSVLFLQTARLVGWQLDDSDV-----VWITGAPK 894

RESULT 8

US-10-449-315-11
; Sequence 11, Application US/10449315
; Publication No. US20030190679A1
; GENERAL INFORMATION:
; APPLICANT: Bertin, John
; TITLE OF INVENTION: NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THEREOF
; FILE REFERENCE: 07334-329001
; CURRENT APPLICATION NUMBER: US/10/449,315
; CURRENT FILING DATE: 2003-05-30
; PRIOR APPLICATION NUMBER: US/09/841,739
; PRIOR FILING DATE: 2001-08-29
; PRIOR APPLICATION NUMBER: US/09/697,089
; PRIOR FILING DATE: 2000-10-26
; PRIOR APPLICATION NUMBER: US/60/161,822
; PRIOR FILING DATE: 1999-10-27
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 11
; LENGTH: 898
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: majority sequence
US-10-449-315-11

Query Match 26.0%; Score 1903; DB 14; Length 898;

Best Local Similarity 50.9%; Pred. No. 1.2e-142;
Matches 484; Conservative 88; Mismatches 214; Indels 164; Gaps 24;

QY 451 EPLVPEVFGNLSVMCVGEAGSGKTVLLKIAFLWASGCCPLNRRFOLVFLVLSSTR 510
DB 1 EQVLVNGVLGALNSVCIVEGAGSGKSVLLQKIAFLWAGKCKALTQFQVFLVLSSTR 60
QY 511 PDEGLASIIICDQLLEKESGVTMCMRNIIQQLKNQVFLFLDDYKEI----CSIPOVIGKL 566
DB 61 ADGGLASIIICDQLLEKESGVTMCMRNIIQQLKNQVFLFLDDYKEI----CSIPOVIGKL 120
QY 567 IQKNHLSRTCLLIATVNRARDIRRYLETILEIKAPFFYNTVCIILKLFSHNMTRLRKPM 626
DB 121 IQENHLSKTCVLVAVTTERARDIROFGALIAEVGAFEDSAVALLREVLIKEIAELRGLL 180
QY 627 VYFGKNSLQIKTKTFLFVAALCA-HWFQYPPDPSPDDVAVPKSYMERSLRNK-----A 680
DB 181 VQIGKSQSIQNLQKTPFLFVAALCAIQWGESFDSFTDVAVPKSFVDDLILKNKHGVA 240
QY 681 TABILKATVSSCGDLALEGVFHKFDFFELDDVAEAGVDEVDLLTTGLLSKFTAQRLRPY 740
DB 241 AADILKATVSSCGDLALEGVFHKFDFFELDDVAEAGVDEVDLLTTGLLSKFTAQRLKPY 300
QY 741 RFLSPAFQBLAGRLIIELDSDROHQDILGLVHLKQINSMPMTVSAYNNFLNYSSLS 800
DB 301 KFLSKAFQBLAGRLIIELDSDROHQDILGLVHLKQINSMPMTVSAYNNFLNYSSLS 360
QY 801 TKAGPKIVSHLLHLVDNKESLENISENDYLLKHQPEISIQMOLLRLGLWQICQAFYSMS 860
DB 361 VEAGRAVSHLAAVVDNKGSLGLSIADYLLKHQESISIQMOLLQGVKNITQAILSAVS 420
QY 861 EHLVLALKTAVQSNNTAACSPVLQ-FLQGRITLIGALNL-QY---PFDDH-PESLSLR 914
DB 421 INLLVLGITAYQSSITVAALSQVLEAFLOQKSLTLGAGNLPDYLDFDHLFESASALD 480
QY 915 SIHFPIRGNTKSPRAHFSVLETCFDKSVQPTIDQDY--ASAFEPNWEERNIAEKEDNVK 972
DB 481 SIKLSIRGGATASRAKAAVLTGGIDKSEAPTIDETIYPASAVSLFENEQEALATLEVTVK 540
QY 973 SYMDMQRASPDLSGTGYWKLSPQYKIP-----CLEVDVNDIDVV 1012

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Db 541 SFSDLNKQAITDLGTGFSASSLQIQIKRCAGVAGSLVLSLSTCKNIYSLEVDASDLTV 600
Qy 1013 GDMLEIL--MTVFS-----ASORIELNLHS-----RGFTESIRPALELSKASVTK----- 1057
Db 601 GEDHUTIVTNLVSIHDLASORLGGTDSLGNKGLIELIRDALELSEASAIKLAAGL 660
Qy 1058 -----CSISKLELSAABQELTLTPSLES-----LE-----VSGTIQSDQDQIFPNL 1098
Db 661 KNLKMWCLISLLELSAAGEGLLIVKLSSEBPCDLEELQVSCCLVAGAVQILAQLHNL 720
Qy 1099 DK--FLCLKELSDVLEGINVPSVTPERPFPNPHMEKLLIQISAYDPSKLVKLQNSPN 1156
Db 721 VKLSILDLSLSELDVLDGNIHVSVPDEFNVLQELTALLQIGADV----- 767
Qy 1157 LHVFLKCNFPDFSGSLMTMLVSKCKLTIKFSDFSFFQAVPPVASLPNPFISIKIILNLSG 1216
Db 768 -----GSLSSL-----VASLEEVISLVILGLEG 791
Qy 1217 QPPDEETSEKFAYI--LGSLSNLEELIPTGD-----GIYRVAKL----- 1254
Db 792 QLTDTETISLIGAFIGLGSLSNLEELIAGGDVSSDGLAFMGVFEVAKLVPPDFSTKEP 851
Qy 1255 -----IIQCCQQLHCLRVLSPPKT-----LNDSVVEIAKVAISGFGQ 1292
Db 852 LPDPALVQQLSQV--LSVLSFLQTLARLVGWQLDDDSV-----VWITGAFK 894

RESULT 9
US-10-029-386-33933
; Sequence 33933, Application US/10029386
; Publication No. US20030194704A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharon G.
; APPLICANT: Rank, David R.
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR C
; TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
; FILE REFERENCE: AEOMICA-X-2
; CURRENT APPLICATION NUMBER: US/10/029,386
; CURRENT FILING DATE: 2001-12-20
; NUMBER OF SEQ ID NOS: 34288
; SOFTWARE: Anomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 33933
; LENGTH: 203
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AC005031.1
; OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 1.6
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.95
; OTHER INFORMATION: SWISSPROT HIT: Q13075, EVALUATION 1.00e-112
US-10-029-386-33933

Query Match 14.2%; Score 1037; DB 14; Length 203;
Best Local Similarity 100.0%; Pred. No. 2.1e-74;
Matches 203; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 389 MAQGEAQWFOBAKNNLEQLRAAYTSASFRHMSLLDISDLATDHLGCDLSIASKHISKP 448
Db 1 MAQGEAQWFOBAKNNLEQLRAAYTSASFRHMSLLDISDLATDHLGCDLSIASKHISKP 60
Qy 449 VOEPLVLEVEGNNLSVMCVGEAGSGTKVLLKTAFLWASGCCPLNRFQVFLSLSS 508
Db 61 VOEPLVLEVEGNNLSVMCVGEAGSGTKVLLKTAFLWASGCCPLNRFQVFLSLSS 120
Qy 509 TRPDEGLASIIICDQLLEKEGVSVMCMRNIIQOLKNQVFLFLDDYKEICSPQVIGKLIQ 568
Db 121 TRPDEGLASIIICDQLLEKEGVSVMCMRNIIQOLKNQVFLFLDDYKEICSPQVIGKLIQ 180
Qy 569 KNHLSRTCLLIATVTRNRDIRR 591
Db 181 KNHLSRTCLLIATVTRNRDIRR 203
```

RESULT 10

US-09-841-739-5
; Sequence 5, Application US/09841739
; Patent No. US20020034784A1
; GENERAL INFORMATION:

; APPLICANT: Bertin, John

; FILE REFERENCE: 07334-329001

; CURRENT APPLICATION NUMBER: US/09/841,739

; CURRENT FILING DATE: 2001-08-29

; PRIOR APPLICATION NUMBER: US 09/697,089

; PRIOR FILING DATE: 2000-10-26

; PRIOR APPLICATION NUMBER: US 60/161,822

; PRIOR FILING DATE: 1999-10-27

; NUMBER OF SEQ ID NOS: 16

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 5

; LENGTH: 1204

; TYPE: PRT

; ORGANISM: Homo sapiens

US-09-841-739-5

Query Match 9.0%; Score 661; DB 9; Length 1204;

Best Local Similarity 23.5%; Pred. No. 4.4e-43;

Matches 268; Conservative 199; Mismatches 427; Indels 245; Gaps 43;

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Qy 341 NCPFLQNMK--SSAEVTPD---LQSRGELCELLETTSENLSDSIAGVIPVEMAQGEAQW 396
Db 216 NYPLFQDLNGQSPBETQNWFFNTSSLIGLFHTQSEGLDD----- 257
Qy 397 FQEAKNLNEQLRAAYTSASFRHMSLLDISDLATDHLGCDLSI---ASKHISKV--- 449
Db 258 -----LAQDLKDLVHTPSFLNF-----YPLGEDIDIIFNKSTFTEPVLWRK 299
Qy 450 -----OEPLVLEVEGNNLSVMCVGEAGSGTKVLLKTAFLWASGCCPLNRFQVFLV 503
Db 300 DQHHHRVEQLTLNGLLQALQSPCIEGSGKGKSTLLQRIAMLGSGKCKALTKEKVFVF 359
Qy 504 LSLSTRPDEGLASIIICDQLLEKEGVSVMCMRNIIQOLKNQVFLFLDDYKEICSPQ-- 561
Db 360 LRLS--RAQGLFETLDCDQLDIPGTTIRKQTFMAMLLKLRQVFLFLDDYNEF--KPQNC 415
Qy 562 -VIGKLIQKNHLSRTCLLIATVTRNRDIRRIVLETLILKAPFPVNTVCILRKLFSHNMT 620
Db 416 PEIETALIKENHRFRKNNVIVTTTTECLRHIRQFGALTAEVGMTEDSAQALIREVLKELA 475
Qy 621 RLRFKFMVYFGKNQSLQIKTQTFLEFVAATCAHFQYFPDPSPDDVAVFKSYMRLSLRNK- 679
Db 476 --EGLLLIQIKSRCLRNLMKTPLFVVITCAIQMGSEBPHSHTQTTLFHTFVDDLKQNKH 533
Qy 680 -----ATAILKATVSSCGELALKGFFSCCFEFDNDLAEAGVDEDEDLTCLMSKFTAQ 734
Db 534 KHKGVAASDFIR-SLDHCGDLALBGFVSHKDFELQDV--SSVNEVDVLLTTGLLCKYTAQ 590
Qy 735 RLRFYPLSPAFOEFAGMELIELDSDROHQDLGLYHLKQINSPMWTVSAYNNFLNY 794
Db 591 RPKPKYKFFHKSFQYETAGRLSLSLTSHBPEVTKNGYIQLQKMWISDITSTYSLLRY 650
Qy 795 V--SSLPSTKAGPKIVSHLLHLVDN-----KESLENISENDYLKHQPEI 837
Db 651 TCGSSVEATRA---VMKHAAVYQHGCLLGLLSIAKRPWRQESLQSVKN----- 696
Qy 838 SLQMLRLGLMQLCPQAYFMSVSEHLLVLAUKTAYQSN--VAACSPFVQLFQGLRTLIG 896
Db 697 TTEQELIKAI---NINSFVECGIHL-----YQESTSKSALSQBFAFQKSLVIN 744
Qy 897 ALNL-QYFPDFHPSLSLLRSHFPIRGNKTSFRAHFSVLETCFDKXSVQPTIDODVASAFE 955
Db 745 SGNIPDYLFD-----FFEHLPNC--ASALDFIKLDFYGG-- 776
Qy 956 PMNEWERNLAEKEDNVKSYMMDQRRASPDLSLSTGYWKLSPKQYKIP----- 1000
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Db 777 AMASWE-----KAAEDTGGIHEEAPETY-IPSRVSLFFNWKQBF 817
QY 1001 CLEVVDNDIVVGQDMLEILMTVFSASQRIELHLSHSGFIESIRPALELSKASVTKCSI 1060
Db 818 TLEVLRDFSKLNKQDIRYLKGFSSATSLRQIKRCAGVAGSLSLVSTCK-NIYSLMV 876
QY 1061 SKLELSAAEQLELLTLPISLESLEVSQTIQSDQIFPN--LDKFLCLKELSDVLEGNINVF 1118
Db 877 EASPLTIEDERHITSVNLKLSIH--DLQNRLPGGTDSLGNLKNLTKLIMDNKM- 932
QY 1119 SVIPEEFPNFHMEKLLIQISAEDPSKLVKLIQNSPNLHVPHLKNFSDPGSLMTMLV 1178
Db 933 -----NEEDAIKLAEBGLKNLKKMCLFHL--THLSDIGEGMDYIV 969
QY 1179 S-----CKKLTIEKFSDFP--QAVPFVA-SLPNFISILKILNLSGQQFPDDETKFAY 1229
Db 970 KSLSEPC-DLEEIQLVSCCLSANAVKILQNLHNLVLSLIDL-SENYLEKDGNEALHE 1027
QY 1230 ILGSLSNLEE--LILPTGDGIYRVAKLIIQOCQQLHCLRLVLSFFKTLNDDSVVEIAKVA 1286
Db 1028 LIDRMNVLEQLTALMLPWGCDVQGSLSLLKHLEVPQVLKGLKNWRLTDI--EIR 1082
QY 1287 ISGGF-----QKLENLKLINSIHKITEGYRNFQALDNMPLQELDISRHTECIKAQA 1340
Db 1083 ILGAFFGNPLKNFQQLNLAGN-RVSSDGLAFMGVFENLKQLVFFDFS---TKEFLPDP 1138
QY 1341 TTVKLSQCVLRPLRLIRLNMLSWLLDADDIALLNVKMERHPQSKVLTILQKWILFPSP 1399
Db 1139 ALVRKLSQVLSKLTFLQEARLVGMQFDDDDLSVITDEKAQ-----MICPWVILKLP 1189

RESULT 12
US-10-449-315-5
; Sequence 5, Application US/10449315
; Publication No. US20030190679A1
; GENERAL INFORMATION:
; APPLICANT: Bertin, John
; TITLE OF INVENTION: NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THERE
; CURRENT APPLICATION NUMBER: US/10/449,315
; CURRENT FILING DATE: 2003-05-30
; PRIOR APPLICATION NUMBER: US /09/841,739
; PRIOR FILING DATE: 2001-08-29
; PRIOR APPLICATION NUMBER: US 09/697,089
; PRIOR FILING DATE: 2000-10-26
; PRIOR APPLICATION NUMBER: US 60/161,822
; PRIOR FILING DATE: 1999-10-27
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 5
; LENGTH: 1204
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-449-315-5

Query Match 9.0%; Score 661; DB 14; Length 1204;
Best Local Similarity 23.5%; Pred.No. 4.4e-43;
Matches 268; Conservative 199; Mismatches 427; Indels 245; Gaps 43;

QY 341 NCPFLQNMK-SSAEVTPD---LQSRGELCELLETTSESLENLSIAVGPITVPERMAQGEAQM 396
Db 216 NYPFLQDLNGQSFEEQNWFNFTSSLIGLPHOTSEGDLDD----- 257
QY 397 FQBAKNLNQLRAAYTSASFRRMSLLDISDLDATDHLGLCDLSI----ASKHISKPV---- 449
Db 258 -----LAQDLKDYHTPSFLNP-----YPLGEDIDIIIFNLKSTFTPEYLMWK 299
QY 450 -----QEPVLPEVFGNLSVMVCVEGAGSKTVLLKKIAFLWAGCCPLNRFOLVFEY 503
Db 300 DQHHHRVEQLTNGLLQALQSPCIIIEGSGKGKSTLLQRIAMLWGSCKKALKTKFKEVFF 359
QY 504 LSLSTRPDEGLASIIICDQLLEKEGSGVTMCMRNI IQLKNOVLFLDDYKEICSPQ-- 561
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Db 360 LRLS--RAQGLFETLCDQLDIPGTIRKQTFMAMLLKLRQRLVFLFLDGYNEP--KPQNC 415
QY 562 -VIGKLIQKNHLSRTCLLIIVRTNRARDIRRYLETILBEIKAPFPVNTVICRLKLPFSHMT 620
Db 416 PEIEALIKENFRKNMVIIVTTTTECLRHIRGFGALTEVGMDETSQAALLREVLIKELA 475
QY 621 RLRFMVYFGNQSLQIKTLPFVAACAHMFQVFPDPSPDDVAVFKSYMERLSLNRK- 679
Db 476 --EGLLQIQKSRCLNLMKTPLFVITCAIQMGSEBFSHTQTTLTFHTFYDLLIKQNKH 533
QY 680 -----ATABIILKATVSSCGELAKGFFSCCFEFDNDLLAEAGVDEDELTWCLMSKPTAQ 734
Db 534 KHKGVAASDFLR-SLDHCGDLALEGVFHKHFDQLQV--SSVNEVDVLLTGLLCKYTAQ 590
QY 735 RLRFYRFLSPAFOEFLAGMELIELLSDSDROHODGLYHLKQINSPMVTSAVNNFLNY 794
Db 591 RPKPKYFTHKSFQBYTAGRLSSLLTSHPEEVTKGNGYLOKQWVSIIDISTVSSILRY 650
QY 795 V--GSLPSTKAGPKIVSHLLHLVDN-----KESLENISENDDYLKHQPEI 837
Db 651 TCGSSVEATRA--VMKILAAVYQHGCLLGLSLAKRPLWRQESLQSVKN----- 696
QY 838 SLQMOLLRLGLWQICQAVFMSVSEHLLVLAKTAYQSNF--VAACSPFVLQFLOGRTILIG 896
Db 697 TTEQILKAI---NINSFVECGIHL-----YQESTSKSALSQEPFAFPQGSLEYIN 744
QY 897 ALNL--QYFFDHPELSLLRSIHFPIRGNKTSIPRAHFSVLETCFDKSOVPTTDQDVASAFE 955
Db 745 SGNIPDYLPD-----FFHELPGN--ASALDFIKLDPYGG-- 776
QY 956 PMNEWERNLAEKEDNVKSYNDMORRAASPDLSGTGYWKLSPKQYKIP----- 1000
Db 777 AMASWE-----KAAEDTGGIHEEAPETY-IPSRVSLFFNWKQBF 817
QY 1001 CLEVVDNDIVVGQDMLEILMTVFSASQRIELHLSHSGFIESIRPALELSKASVTKCSI 1060
Db 818 TLEVLRDFSKLNKQDIRYLKGFSSATSLRQIKRCAGVAGSLSLVSTCK-NIYSLMV 876
QY 1061 SKLELSAAEQLELLTLPISLESLEVSQTIQSDQIFPN--LDKFLCLKELSDVLEGNINVF 1118
Db 877 EASPLTIEDERHITSVNLKLSIH--DLQNRLPGGTDSLGNLKNLTKLIMDNKM- 932
QY 1119 SVIPEEFPNFHMEKLLIQISAEDPSKLVKLIQNSPNLHVPHLKNFSDPGSLMTMLV 1178
Db 933 -----NEEDAIKLAEBGLKNLKKMCLFHL--THLSDIGEGMDYIV 969
QY 1179 S-----CKKLTIEKFSDFP--QAVPFVA-SLPNFISILKILNLSGQQFPDDETKFAY 1229
Db 970 KSLSEPC-DLEEIQLVSCCLSANAVKILQNLHNLVLSLIDL-SENYLEKDGNEALHE 1027
QY 1230 ILGSLSNLEE--LILPTGDGIYRVAKLIIQOCQQLHCLRLVLSFFKTLNDDSVVEIAKVA 1286
Db 1028 LIDRMNVLEQLTALMLPWGCDVQGSLSLLKHLEVPQVLKGLKNWRLTDI--EIR 1082
QY 1287 ISGGF-----QKLENLKLINSIHKITEGYRNFQALDNMPLQELDISRHTECIKAQA 1340
Db 1083 ILGAFFGNPLKNFQQLNLAGN-RVSSDGLAFMGVFENLKQLVFFDFS---TKEFLPDP 1138
QY 1341 TTVKLSQCVLRPLRLIRLNMLSWLLDADDIALLNVKMERHPQSKVLTILQKWILFPSP 1399
Db 1139 ALVRKLSQVLSKLTFLQEARLVGMQFDDDDLSVITDEKAQ-----MICPWVILKLP 1189

RESULT 12
US-10-156-733-2
; Sequence 2, Application US/10156733
; Publication No. US20030099969A1
; GENERAL INFORMATION:
; APPLICANT: Alnemri, Emad S.
; TITLE OF INVENTION: IPAF, AN ICE-PROTEASE ACTIVATING
; TITLE OF INVENTION: FACTOR
; FILE REFERENCE: 480140.477
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Db 726 IEDERHITSVTNKLTSIH---DLQNRPLPGGLTDSLGNLKNLTKLIMONIKM-----775
QY 1125 FNFHMEKLLIQISAEDPSKLVKLIQNSPNLHVFLKCNFSDFGSLMTMLVS-----1179
Db 776 -----NEEDAIAELGLKNLKNWCLFHL--THLSDLGEGMDYIVKLSSE 818
QY 1180 -CKKLTETKFSDFP--QAVPFA-SLPNFIKILNLGSGQFPDEETSEKPAYILGSL 1235
Db 819 PC-DLEEIQLVSCCLSANAVKILAQNLHNLVKLSILD--SENYLEKDGNEALHELIDRMN 876
QY 1236 NLEE---LILPTGDCIYRVAKLIIQCCQQLHCLRVLSFFKTLNDDSVVEIAKVAISGGP- 1291
Db 877 VLEQLTALMLPGWCDVQGSLSLLKHLSEVPQLVGLGNLWRLTDT-----BIRILGAFP 931
QY 1292 -----QKLENKLSINHKITEGYRNFQALDNMPNLQBLDISRHFTCEIKAAQATTVKSL 1346
Db 932 GKNPLKNFOQLNLAGN-RVSSDGLAFMGVFNELKQLVFFDFS---TKFELPDPALVRKL 987
QY 1347 SQCVLRPLRLIRLNLMSWLLDADDTALL 1374
Db 988 SQVLSKLTFLQEARLVGMQFDDDDLSVI 1015

RESULT 15
US-10-449-315-2
; Sequence 2, Application US/10449315
; Publication No. US20030190679A1
; GENERAL INFORMATION:
; APPLICANT: Bertin, John
; TITLE OF INVENTION: NOVEL MOLECULES OF THE CARD-RELATED PROTEIN FAMILY AND USES THERE
; FILE REFERENCE: 07334-329001
; CURRENT APPLICATION NUMBER: US/10/449,315
; PRIOR FILING DATE: 2003-05-30
; PRIOR APPLICATION NUMBER: US/09/841,739
; PRIOR FILING DATE: 2001-08-29
; PRIOR APPLICATION NUMBER: US 09/697,089
; PRIOR FILING DATE: 2000-10-26
; PRIOR APPLICATION NUMBER: US 60/161,822
; PRIOR FILING DATE: 1999-10-27
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 1024
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-449-315-2

Query Match 8.9%; Score 653; DB 14; Length 1024;
Best Local Similarity 23.7%; Pred. No. 1.5e-42;
Matches 263; Conservative 197; Mismatches 410; Indels 238; Gaps 42;

QY 344 FLQNMKS-SAEVTPDLQSRGECLELLETTSESNLEDSTIAGVPIVPEMAQGEAQWFOEAKN 402
Db 69 FLKSLKEMNYPFLQDLNGQ-----SLFHQTSEGLDD-----100
QY 403 LNEQLRAAYTSASFRHMSLLDISSDLATDHLIGCDLSI---ASKHISKPV-----449
Db 101 LAQDLKDLVHTSPFLNF-----YPIGEDIDIIIFNLKSTFTFEPVLWRKQDHHHR 148
QY 450 QBPLVLPVEFGNLSVMCEGABAGSKTVLLKKIAFLNAGSCCLLNRFQVLYLSLSST 509
Db 149 VEQLTLNGLLQALQSPCIIIEGSGKSTLLQRIAMNGSGKCKALTREKFFVFLRLS--206
QY 510 RDEGLASIIICQLLEKGSVTMCMWRNIIQQLKNQVLPFLDDYKEISIPQ---VIGKL 566
Db 207 RAQGGFLFETCLDQLLDIPGTIRKQTFMAMLLKLRQLRVFLPDGYNEF--KPQNCPEIEAL 264
QY 567 IQKNHLSRTCLLIJAVTNRADIRVLETILIKAFPPYNTVCILRKLFSHNWTLRKFM 626
Db 265 IKENHRFNKMVIIVTTTTECLRHIFQFALTAEGVGMNTEDSAQALLIREVLIKELA--EGLL 322
QY 627 VYFGKNQSLQIKQKTPLFVAALCAHWFPDFPDSFDDVAVFKSYMERSLRNK-----A 680
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Db 323 LQIQSRCLRLNMLKPTLFVVIITCAIQMGESEFHSHTQTTLFTFYDLLLQKNKHKGVA 382
QY 681 TAEILKATVSSCGELALGFFSCCFEFNDDDLAEAGVDEDEDLTCLMSKFTAQRLPFY 740
Db 383 ASDFR-SLDHCDLALGEGVFSHKDFELODV--SSVEDVLLTTGLLCKYTAORFKPKY 439
QY 741 RFLSPAFOEFLAGMLRIELLDSDROHODGLYHLKQINSPMTVSYANNFLNYV--SSL 798
Db 440 KPFHKSFOEYTAGRLSLLTSHPEEVTKGNGYLQKMWISIDITSTYSLLRYTCGSSV 499
QY 799 PSTKAGPKIVSHLLHLVDN-----KESLENISENDDYLKHQPEISLQML 843
Db 500 EATRA---VMKHAAYVQHGLLGLSIAKRLPMWRQESLQSVKN-----TTEQEI 545
QY 844 LRGLMQICPQAVFSVMVSHLLVLALKTAYQSN--VAQSPFVLQFLQGRTLTLGALNL-Q 901
Db 546 LKAI-----NINSVFCGIHL-----YQESTSKALSQEFQKSLYINSQNI 593
QY 902 YFFDHPESLSLRSIHFPTRGNKTSIPRAHFSVLETCFDKSOVPTIDQDYASAFEPNME 961
Db 594 YLFO-----FFEHLNPN--ASALDFIKLDFYGG--AMASWE 625
QY 962 RNLAEDNVKSYMDMQRASPDLTGTGKWSPKQYKIP-----CLEVDV 1006
Db 626 -----KAAEDTGGIHMEEAPETY-IPSRVSLFFNWKQBFRTLEVTL 666
QY 1007 NDIDVVGQDMLIEMTVFSASQRIELHNLHSGFTIESIRPALELSKASVTKCSISKLELS 1066
Db 667 RDFSINKQDITYLGKIFSSATSLRQIKRCAGVAGSLVLSTCK-NIYSLMVEASPLT 725
QY 1067 AAEQELLTLPLSLESLEVSIGTISQDQIFPN--LDKFLCLKELSDLEGNINVSFVPIEE 1124
Db 726 IEDERHITSVTNKLTSIH---DLQNRPLPGGLTDSLGNLKNLTKLIMDNIKM-----775
QY 1125 FNFHMEKLLIQISAEDPSKLVKLIQNSPNLHVFLKCNFSDFGSLMTMLVS-----1179
Db 776 -----NEEDAIAELGLKNLKNWCLFHL--THLSDLGEGMDYIVKLSSE 818
QY 1180 -CKKLTETKFSDFP--QAVPFA-SLPNFIKILNLGSGQFPDEETSEKPAYILGSL 1235
Db 819 PC-DLEEIQLVSCCLSANAVKILAQNLHNLVKLSILD--SENYLEKDGNEALHELIDRMN 876
QY 1236 NLEE---LILPTGDCIYRVAKLIIQCCQQLHCLRVLSFFKTLNDDSVVEIAKVAISGGP- 1291
Db 877 VLEQLTALMLPGWCDVQGSLSLLKHLSEVPQLVGLGNLWRLTDT-----BIRILGAFP 931
QY 1292 -----QKLENKLSINHKITEGYRNFQALDNMPNLQBLDISRHFTCEIKAAQATTVKSL 1346
Db 932 GKNPLKNFOQLNLAGN-RVSSDGLAFMGVFNELKQLVFFDFS---TKFELPDPALVRKL 987
QY 1347 SQCVLRPLRLIRLNLMSWLLDADDTALL 1374
Db 988 SQVLSKLTFLQEARLVGMQFDDDDLSVI 1015
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Job time : 106 secs

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